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EXPERIMENTAL DESIGNS

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PREFACE

Work on this book was started when both of us were members of the staff of Iowa State College. At that time requests were received rather frequently from research workers. Some wanted advice on the conduct of a specific experiment: others, who had decided to use one of the more complex designs that have been discovered in recent years, asked for a plan or layout that could be followed during the experimental operations. Although the logical principles governing the subject of experimentation are admirably expounded in Fisher's book *The Design of Experiments*, these requests indicated a need for a different type of book, one which would describe in some detail the most useful of the designs that have been developed, with accompanying plans and an account of the experimental situations for which each design is most suitable. Such a book is directed at the experimenter and is intended to serve as a handbook which is consulted when a new experiment is under consideration.

Mainly on account of the war, slow progress was made. In 1944 we completed a mimeographed draft of which several hundred copies were distributed. Many helpful suggestions were made by readers. Of these, the most frequent was to the effect that we should include more material dealing with the statistical analysis of the results. In the mimeographed draft, our practice was to give references to worked examples of the analysis in cases where they could be found in the literature, and to present examples for only those designs whose analysis was not available. To this it was objected that many research workers did not have easy access to our references.

This suggestion raised a difficult issue. To present a self-contained account of the analysis of variance in all its ramifications would make the book, it seemed to us, unwarrantably long and expensive. Consequently we have continued to assume that the reader has some knowledge of the principles of analysis of variance and of the computational methods involved. We have included a brief review of the basic theory and an extensive set of worked examples of the analysis for both common and less common types of design. Although strenuous efforts were made to obtain a selection of examples from diverse fields of research, a preponderance from biology, and more particularly from

agriculture, was dictated by our own experience in those areas. On several occasions we rejected an example which would have made an attractive addition to the scope because we did not feel sufficiently familiar with the conditions of experimentation to give a realistic account of the problems encountered.

Since courses of lectures on the design of experiments are being introduced in many colleges and universities, some teachers may be interested in the potentialities of the book as a textbook. Several comments are prompted by our own experience in this connection. First, it will often be necessary for the teacher to provide a more systematic development of the analysis of variance than is given here. Second, interest in such a course is greatly enhanced by examples from an environment with which the listeners are familiar, and especially by examples of experiments that have been conducted by some of the listeners. Thus the teacher is well advised to build the course around his own examples, using those in this book mainly as supplementary material for the students. Finally, a selective use of the book is in order, because it contains much more material than can be covered in a typical one-quarter course, and because some difficult topics have been dealt with in the early parts of the book.

We wish to express our gratitude to the many staff members at Iowa State College and North Carolina State College who helped us by putting experimental data at our disposal, by providing painstaking descriptions of their experimental techniques, or by lending their assistance in the preparation of the manuscript and plans. For similar kindnesses in connection with certain of the examples we are indebted to G. F. Potter, D. Y. Solandt, D. B. DeLury and J. Hunter, F. M. Wadley and C. F. Rainwater, and the Wailuku Sugar Company, Honolulu. Some theoretical results used in Chapter 14 were developed from research conducted at Raleigh under a contract with the Office of Naval Research. Finally, our thanks go to George W. Snedecor, who participated in the original plans for this book and made a careful reading of the first draft.

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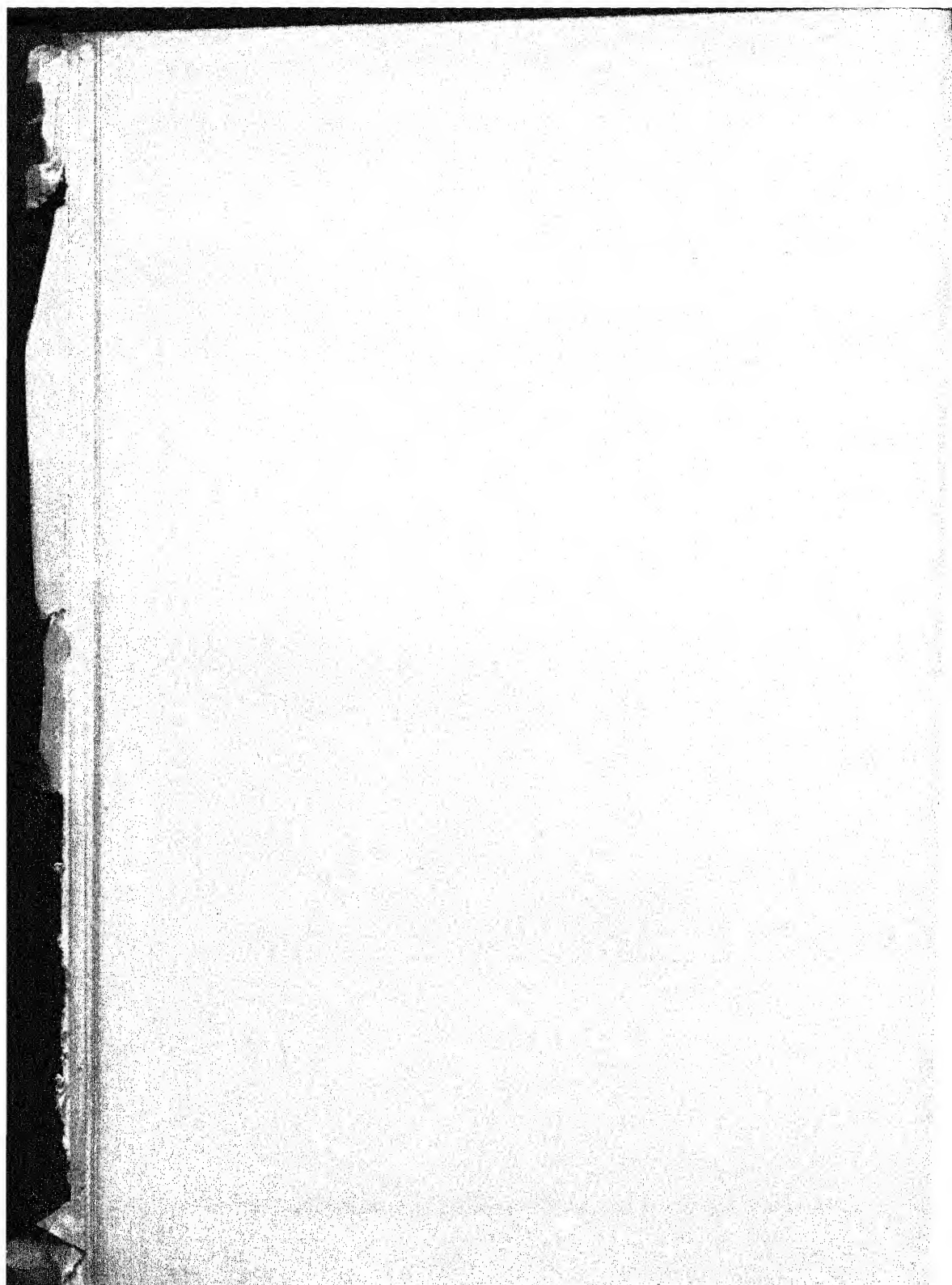
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CHAPTER 1

INTRODUCTION

1.1 The Contribution of Statistics to Experimentation

1.11 The Problem of Interpretation. Since statisticians do not usually perform experiments, their claim to attention when they write on this subject requires some explanation. It is true that on many important aspects of experimentation the statistician has no expert knowledge. Nevertheless, in recent years, research workers have turned increasingly to statisticians for help both in planning their experiments and in drawing conclusions from the results. That this has happened is convincing evidence that statistics has something to contribute.

At first, requests for assistance were nearly always concerned with the interpretation of the results. It is a common characteristic of experiments in widely diverse fields of research that, when they are repeated, the effects of the experimental treatments vary from trial to trial. This variation introduces a degree of uncertainty into any conclusions that are drawn from the results. Even after a number of repetitions, or *replications* as they are called, the investigator still does not know by how much his results would be changed if the experiment were repeated further under the same conditions. Successive trials may be so discrepant in their results that it is doubtful which of two treatments would turn out better in the long run.

As an illustration of this variation, data are given in table 1.1 from a simple experiment. The data are the times (minus 2 minutes) required to perform a routine statistical calculation, that of finding the sum of squares of 27 observations. Each sum of squares was computed separately by the same person on both of two standard computing machines. In all, 10 different sums of squares were worked, making 10 trials or replications of the experiment. It will be noted that the differences in speed range from 17 seconds in favor of machine *B* to 2 seconds in favor of machine *A*. Some experimenters may comment that the results of this experiment are remarkably well behaved, and exhibit nothing like the variation with which they have to contend. The results will, however, serve our purpose.

TABLE 1.1 TIME IN SECONDS (MINUS 2 MINUTES) REQUIRED FOR COMPUTING SUMS OF SQUARES

Replication	Machine		Difference ($A - B$)
	A	B	
1	30	14	16
2	21	21	0
3	22	5	17
4	22	13	9
5	18	13	5
6	29	17	12
7	16	7	9
8	12	14	-2
9	23	8	15
10	23	24	-1
Means	21.6	13.6	8.0

The object of the experiment is, of course, to compare the speeds of the two machines for this calculation. More specifically, two objects might be stated. The first is to answer the question: is there any difference in speed? or, to put it another way, to test the hypothesis that there is no difference in speed. The second object, which is related to the first, is to estimate the size of the difference in speed. Almost all experiments are carried out for one or both of these purposes—the testing of hypotheses and the estimation of differences in the effects of different treatments.

As regards the test of the hypothesis that there is no difference in speed, we might report, as relevant evidence, that B proved faster 7 times out of 10, A twice, while once there was a tie. For the problem of estimation we might report that the average difference in speed in the experiment was 8 seconds in favor of B . These purely descriptive statements do not carry us very far. Their weakness is that they supply no information about the reliability of the figures presented. For example, have we any confidence that, if the experiment were continued for another set of 10 trials, the advantage at the end would still be close to 8 seconds in favor of B ?

Because of the deficiencies in the descriptive approach, experimenters have adopted a different point of view in the summarization of their results. They tend to reason as follows. Suppose that it were feasible to continue the experiment indefinitely under the same conditions. The average difference in speed between the two machines would presumably settle down to some fixed value. This value, which will be independent of the size of the experiment that was actually carried out, may reason-

ably be called the *true* difference between A and B . From this point of view, the problem of summarizing the results may be restated in the question: what can we say about the true difference between A and B ? This is a problem in *induction* from the part to the whole, or in statistical language, from the sample to the population. A solution to this problem has been developed by means of the theory of statistics. It is this solution that constitutes the principal contribution of statistics to the interpretation of the results.

1.12 Statistical Inference. Obviously, it cannot be expected that the solution will provide the exact value of the unknown true difference. As a less ambitious goal we might hope to be able to find 2 limits within which the exact value is certain to lie, but even this cannot quite be attained. What can be done is that for any chosen probability, say .95, two limits are found such that the probability that they enclose the true difference is .95. In other words, limits can be found that are almost certain to enclose the true difference, where the degree of certainty, as measured by the probability, can be chosen by the experimenter. Since we wish to focus attention on the type of inferential statement that can be made rather than on the method of calculating the limits, the computations will not be discussed at present. For the example in table 1.1 they will be found in section 4.42. When the probability is .95, the limits for the true difference in speed between the 2 machines turn out to be 3.3 and 12.7 seconds in favor of machine B . A statement that B is faster by an amount that lies between 3.3 and 12.7 seconds has a 1 in 20 chance of being wrong. If the degree of certainty is decreased by lowering the probability to .8, the limits are narrowed to 5.1 and 10.9 seconds. If the probability is raised to .99, the limits become 1.1 and 14.9 seconds, and as the probability is brought closer to certainty, the limits steadily become farther apart. The limits are called confidence limits, and the probabilities are called confidence probabilities.*

These probabilities are not merely academic abstractions: they can be subjected to at least a rough experimental verification. For verification, we need a situation where the true difference between the effects of two treatments is known. In toxicology, for instance, this situation can sometimes be obtained by diluting a standard poison to a known extent. The dilution is sent to the laboratory, labelled as an "unknown" poison. By experiments on animals, confidence limits are found for the

* Fisher (1.2) has developed statistical inference in terms of *fiducial* limits. The two concepts, fiducial and confidence limits, have different logical backgrounds, although in all simple applications to controlled experiments the actual values of the fiducial and confidence limits are identical. For a discussion, see Kendall (1.1).

amount of the unknown that has the same toxicity as a given amount of the standard. Since the persons originating the experiment know the true value for this amount, they can verify whether the statement that the true amount lies within the limits is correct. The practical difficulty with this type of check is the labor required. A large number of experiments would be needed to verify whether about 95% of the statements made with confidence probability .95 were in fact correct, and about 5% were wrong.

As we have seen, the statistical solution to the problem of estimation consists of a statement that the true difference lies between certain limits, plus a probability that the statement is correct. It is of interest to consider whether this type of information is sufficiently precise to permit decisions of practical importance to be made. Although a thorough discussion of this question would be rather lengthy, inferences of this type often permit definite action to be taken with confidence that the action will be fruitful. When they fail to decide the point at issue, the reason is nearly always that the data obtained are insufficient. For illustration, suppose that it is desired to discover whether the application of a dressing of some fertilizer to a crop will be profitable. The cost of the fertilizer is such that its application will be profitable only if it increases yields on the average by 2 or more bushels. A series of experiments is carried out in order to estimate the true average response to the fertilizer. If the 95% confidence limits for the increase due to the fertilizer are 4 and 11 bushels, its use can be adopted with a good deal of assurance that it will be profitable. Similarly, if the 95% confidence limits are -5 and 1 bushels, a decision not to use the fertilizer follows. A case where there is uncertainty occurs when the limits are 0 and 5 bushels. Here it is likely that there will be either a small gain or a small loss, but no recommendation can be made without considerable risk of its being wrong. If it is important to make the correct decision, further experiments must be conducted in order to narrow the distance between the confidence limits.

Thus far we have considered the problem of estimating the true difference between the effects of two treatments. In testing hypotheses, we are interested in the supposition that the true difference has some specified value, most commonly zero. As in the case of estimation, difficulty arises because of the variability that is typical of experimental data. As a result of this variability, the data are never exactly in agreement with the hypothesis, and the problem is to decide whether the discrepancy between the data and the hypothesis is to be ascribed to these variations or to the fact that the hypothesis is not true. The contribution of statistics is the operation known as a *test of significance*.

Essentially, this is a rule for deciding, from examination of the data, whether or not to reject the hypothesis. Such rules are made to satisfy two conditions that are obviously desirable: (i) hypotheses that are true shall be rejected only very occasionally, and the probability of rejection may be chosen by the experimenter; (ii) hypotheses that are false shall be rejected as often as possible.

This technique enables the experimenter to test his hypotheses about the action of the treatments, with the assurance that there is little risk of erroneously rejecting a hypothesis that happens to be correct. Probabilities of .05 and .01 are most commonly used for this risk, and in these cases the tests are said to be made at the 5 and 1% significance levels respectively. These levels are just useful conventions, and a lower probability may be used if the consequences of an erroneous rejection of the hypothesis are very serious. It should be remembered, however, that in lowering this probability value we automatically diminish the chance of rejecting a hypothesis that is false.

A useful property of a test of significance is that it exerts a sobering influence on the type of experimenter who jumps to conclusions on scanty data, and who might otherwise try to make everyone excited about some sensational treatment effect that can well be ascribed to the ordinary variation in his experiment. On the whole, however, tests of significance are less frequently useful in experimental work than confidence limits. In many experiments it seems obvious that the different treatments must have produced some difference, however small, in effect. Thus the hypothesis that there is *no* difference is unrealistic: the real problem is to obtain estimates of the sizes of the differences.

The construction of confidence limits may add something to the interpretation of a test of significance. For instance, suppose that the speeds of the 2 machines in the example had not been significantly different. It is a commonplace that this result would not prove that the 2 machines were identical in speed. However, if the 95% confidence limits for the difference in speed were -2 and $+4$ seconds, we might argue that a true difference of 4 seconds, even if it existed, would be of no practical significance. Consequently, it could be said that for all practical purposes the 2 machines are identical in speed. This is much more positive and useful than the mere statement that the difference in speeds was not statistically significant. If, on the other hand, the confidence limits were -30 and $+32$ seconds, there is no justification for the conclusion that the machines can be regarded as equivalent. All that we have learned is that the data are not sufficiently accurate to show whether there is a difference in speed that is of practical importance.

To summarize: variability in results is typical in many branches of

experimentation. Because of this, the problem of drawing conclusions from the results is a problem in induction from the sample to the population. The statistical theories of estimation and of testing hypotheses provide solutions to this problem in the form of definite statements that have a known and controllable probability of being correct. These statements are specific enough to be useful in deciding whether action can be taken on the basis of the results.

1.13 The Function of Randomization. As would be expected, the type of statistical inference that can be made from a body of data depends on the nature of the data. It is easy to conduct an experiment in such a way that no useful inferences can be made, and many of the experiments brought to the statistician, particularly in earlier years, have been of this type. To take a simple example, suppose that in the comparison of the calculating machines each sum of squares had been computed first on machine *A* and then on machine *B*. Now it is quite possible that increased familiarity with the data will enable the second computation to be done faster than the first. The advantage is unlikely to be great in such an easy calculation; it could be so if the computation were more difficult. (In the experiment this advantage was estimated at about 4 seconds, though the confidence limits for the true advantage are rather far apart.)

If the experiment is conducted in this way, the observed difference in speed ($B - A$) is an estimate of the true difference, plus the unknown difference in speed between a second calculation and a first. Confidence limits set up by statistical techniques apply not to the true difference between *B* and *A* but to the true difference plus this unknown advantage. Consequently, the limits tell us nothing definite about the true difference between *B* and *A*. The interpretation of tests of significance also becomes confused. If *A* is found significantly faster than *B* by ordinary statistical tests, we can be confident that there is a real difference in favor of *A*, since *A* was handicapped in the course of the experiment. But if *B* is found significantly faster than *A*, we do not know what to conclude. In this case we are dealing with a bias whose nature can be anticipated before the experiment has started. In other experiments where less is known about the type of variability that is present, similar biases that are quite unexpected can occur from some apparently innocuous rule about the way in which different treatments are handled.

In order to avoid these biases we need some means of insuring that a treatment will not be continually favored or handicapped in successive replications by some extraneous source of variation, known or unknown. This is done by the device known as *randomization*, due to Fisher. In-

It cannot
be done in the
randomization

stead of performing every calculation first on machine *A*, we apply the principle of randomization by tossing a coin to determine whether *A* or *B* shall be used first in any trial. The decision is made independently in each trial. The effect is that in any trial each machine has an equal chance of being tested under the more favorable conditions. Of course, the result of any specific randomization may favor one or the other treatment. But this happens only to an extent that is allowed for in the calculations that are used for tests of significance and confidence limits. This important result has been illustrated in detail by Fisher (1.2), who has shown how tests of significance and confidence limits can be constructed, using only the fact that randomization has been properly applied in the experiment. Randomization is one of the few characteristics of modern experimental design that appears to be really modern. One can find experiments made 100 or 150 years ago that embody the principles that are now regarded as sound, with the conspicuous exception of randomization.

The occasions on which randomization is required vary with the type of experiment and must be left to the judgment of the experimenter. One occasion arises when the treatments are allotted to the experimental material. Suppose that the effects of different diets on the heights and weights of children are to be ascertained. Since different children grow at different rates, a treatment that happens to be assigned to a group of fast-growing youngsters is favored. Consequently, we allot the diets in any replication at random to the children who are to receive them, with a new random allotment in each replication. Similarly, if 4 different oven temperatures for the cooking of roasts are under comparison, the 4 temperatures are assigned at random to the 4 roasts which form the material for any replication. Sometimes this is the only randomization required, but frequently other operations that are carried out in the course of the experiment are also potential sources of bias. With a repetitious operation the order of events may be important, either because a learning process is involved which tends to make later operations better than the earlier ones, or because fatigue tends in the opposite direction. Systematic biases may be guarded against by randomizing the order in which the operation is performed on the different treatments in a replication. In other cases the equipment that is used introduces variation. For example, if 4 ovens are available to cook the 4 roasts, we would not always use the same oven for the same temperature, in case biases should be introduced because of systematic differences among the ovens. Instead, the temperatures could be assigned at random to the ovens in each replication. Thus we have *two* randomizations, one to assign the temperatures to the roasts and one to assign them to the

ovens. If, however, we decide, before randomizing, which roast is to go in which oven, the two randomizations can be reduced to one. This method can always be used, if convenient, to cut down the number of randomizations that must be made.

Randomization is somewhat analogous to insurance, in that it is a precaution against disturbances that may or may not occur and that may or may not be serious if they do occur. It is generally advisable to take the trouble to randomize even when it is not expected that there will be any serious bias from failure to randomize. The experimenter is thus protected against unusual events that upset his expectations. Of course in experiments where a great number of physical operations are involved, the application of randomization to every operation becomes time consuming, and the experimenter may use his judgment in omitting randomization where there is real knowledge that the results will not be vitiated. It should be realized, however, that failure to randomize at any stage may introduce bias unless either the variation introduced in that stage is negligible, or the experiment effectively randomizes itself.

1.14 Restricted Randomization. We now reconsider the randomization proposed for the experiment on computing machines, in order to discuss a criticism that presumably has occurred to the reader. When this experiment is being planned, we possess the knowledge that there is some advantage to the machine on which the second calculation of any sum of squares is made. In the light of this knowledge, it seems wise to make sure that each machine receives the advantage in five of the 10 replications, rather than to leave this to be decided by the tossing of coins. In fact, if the decision is made by tossing coins, the chances are only about 1 in 4 that each machine will be used first in five replications; they are also about 1 in 4 that one machine will be used first in 7 replications. The criticism is that the randomization is unlikely to give as accurate a comparison as the proposed alternative. The criticism is sound, and the experiment was actually planned so that each machine would be used first in half the replications.

Randomization was also applied in the design actually used, though it is a different type from that originally proposed, being subject to the restriction that each machine must appear first exactly five times. The process of randomizing therefore consisted in choosing five numbers at random from the numbers 1 to 10; these turned out to be 1, 3, 6, 8, and 9. Accordingly, machine A was used first in replications 1, 3, 6, 8, and 9. This is one of the cases where it is difficult to see why failure to randomize could have led to any serious danger of bias. It probably would have been quite satisfactory to use machine A first in the first 5 replicates, and

machine *B* first in the remainder. Nevertheless, it *might* happen that the advantage in the second calculation of a sum of squares would diminish in later replications; and, to save the trouble of trying to guess whether biases of this type are likely, the randomization was used as a precaution.

The example represents a fairly common situation, in which knowledge of the variation that will affect the results is sufficiently detailed so that the simplest type of randomization is objectionable on the grounds of accuracy. Most of the designs in this book were constructed for this situation. By suitable restrictions on the randomization, these designs enable the experimenter to utilize any knowledge that will increase the precision of the experiment. In addition, each design allows sufficient randomization to prevent biases from sources of variation about which knowledge is less certain. For each of the principal designs described, the appropriate method of randomization is also presented.

One further point should be mentioned. We have seen that the experiment on computing speeds could have been carried out either by means of the simple randomization originally proposed or by means of the restricted randomization which ensures that each machine is used first five times. Both methods provide a valid test of significance and valid confidence limits. However, the calculations made for the test of significance and for the confidence limits are different in the 2 cases. The first experiment is of the type known as randomized blocks, and calculations are made as described in section 4.23; the second experiment is of the "cross-over" type, for which the calculations are given in section 4.42. This is an instance of the general rule that the way in which the experiment is conducted determines not only whether inferences can be made, but also the calculations required to make them. The experimenter must make sure that the calculations which he uses are appropriate for the experiment.

1.2 Initial Steps in the Planning of Experiments

1.21 Importance of the Initial Steps. It has been mentioned that statisticians are often asked for advice in making inferences from the results of experiments. Since the inferences that can be made depend on the way in which the experiment was carried out, the statistician should request a detailed description of the experiment and its objectives. It may then become evident that no inferences can be made or that those which can be made do not answer the questions to which the experimenter had hoped to find answers. In these unhappy circumstances, about all that can be done is to indicate, if possible, how to avoid this outcome in future

*do not include
the experiment
which is not
the object of
the experiment*

experiments. Consequently, it has come to be realized that the time to think about statistical inference, or to seek advice, is when the experiment is being planned.

Participation in the initial stages of experiments in different areas of research leads to a strong conviction that too little time and effort is put into the planning of experiments. The statistician who expects that his contribution to the planning will involve some technical matter in statistical theory finds repeatedly that he makes a much more valuable contribution simply by getting the investigator to explain clearly why he is doing the experiment, to justify the experimental treatments whose effects he proposes to compare, and to defend his claim that the completed experiment will enable its objectives to be realized. For this reason the remainder of this chapter is devoted to some elementary comments on the subject of planning. These comments are offered with diffidence, because they concern questions on which the statistician has, or should have, no special authority, and because some of the advice is so trite that it would be unnecessary if it were not so often overlooked.

It is a good practice to make a written draft of the proposals for any experiment. This draft will in general have three parts: (i) a statement of the objectives; (ii) a description of the experiment, covering such matters as the experimental treatments, the size of the experiment, and the experimental material; and (iii) an outline of the method of analysis of the results.

1.22 The Statement of Objectives. This statement may be in the form of the questions to be answered, the hypotheses to be tested, or the effects to be estimated. The aim should be to make the statement lucid and specific. The most common faults are vagueness and excessive ambition, in the sense that a twenty-year research program would be required in order to realize the stated objectives. Often it is advisable to classify the objectives as major and minor, because some types of experiment give high precision for certain treatment comparisons but low precision for others. When the experiment represents a cooperation among people of different interests, this classification is particularly helpful in that it makes clear which objectives take priority and helps to avoid an unhappy compromise that is adopted in the hope of pleasing everybody.

The statement should include an account of the area over which generalizations are to be made, or in other words, of the population about which it is hoped to make inferences. If an experiment is to be conducted on persons suffering from some disease, are the results presumed to apply to the patients in a specific hospital, to patients in all hospitals,

or to all sufferers whether in hospital or not? This sort of question is crucial in applied research, where often the experimenter has some definite population in mind to which he would like to apply the results. It is obvious that worth-while inferences about an extensive population cannot be made from a single experiment. For instance, inferences made from the experiment on the two computing machines are restricted to the person who made the calculations and to the type of calculation made. There is no guarantee that the results would be the same for a different type of calculation or with different operators. Consequently, if the object were to find out which machine is the speedier in general use in a computing laboratory, the experiment that was made has only scratched the surface of the problem.

1.23 The Experimental Treatments. We have used the term *treatments* to denote the different procedures whose effects are to be measured and compared. In the selection of treatments it is important to define clearly each treatment and to understand the role that each treatment will play in reaching the objectives of the experiment. Some issues that arise in particular cases are as follows.

1. Confusion is sometimes caused by failure to distinguish whether the object is merely to "spot the winner" among the different treatments, or whether in addition it is desired to find some clues as to why the treatments behave as they do. A good example is the experiment which demonstrates, or so we are informed, that each of the three treatments, whiskey and water, gin and water, and rum and water, taken orally in sufficient quantities, produces some degree of intoxication. By itself the experiment provides no information as to whether the intoxication is due to the water, to the other ingredient, or to the fact that the two are mixed. A more extensive experiment with additional treatments would be necessary to throw some light on this question. Although there are occasions when it is sufficient simply to discover which is the best of the treatments, experience suggests that even in strictly applied research progress is faster if experiments supply fundamental knowledge. Similarly, the criticism that a certain treatment should be omitted because it would not be used in practice is valid if the aim is to find the best of the "practical" treatments, but it is not valid if the "impractical" treatment will supply some needed information about the behavior of other treatments in the experiment.

2. The specification of the treatments may raise difficult questions about the conditions under which the treatments are to be compared. Suppose that the object is to find out the effect of an application of sulphate of ammonia on the yield of some crop. It is well known that

this effect is likely to depend on the amounts of other important plant nutrients that are available to the crop. Consequently, it must be decided whether to add these nutrients so that they will be in abundant supply on all plots, or to test the nitrogen on the soil as it happens to be. The decision must be guided by the objectives of the experiment, and more particularly by the type of population to which inferences are to apply. Sometimes it is advisable to test the nitrogen in both the presence and the absence of the other nutrients. One result of this decision will be that the experiment becomes also a test of the other nutrients, even though this may not have been part of the original plan. Experiments of this type, where one factor is tested over different levels of another factor, are known as *factorial experiments* and have come to play a prominent part in experimentation. They are discussed more fully in chapter 5.

3. In some cases it becomes apparent that the treatments that can be tested in practice are not those that we would like to test. The following example is typical of many fields where human relations are involved. Suppose that it is desired to compare two methods *A* and *B* of teaching a foreign language, and that we have defined each method clearly and have decided how to measure the success achieved by each method. However, some of the teachers who are to participate in the test already use a method similar to method *A* and have strong beliefs that method *B* is inferior, whereas others already follow method *B* and have no use for method *A*. If the teachers are divided at random into the groups that are to teach the two methods, we may expect that some teachers will be assigned to teach a method in which they have no faith, and we may decide that this is not the kind of comparison that we had in mind. On the other hand, if we forsake randomization and allow teachers to use the method that they like, differences attributed to the methods may in fact be due to differences in the ability of the two groups of teachers. Given enough resources, the situation can be more thoroughly explored by having each method taught by those who like it, those who do not, and those who are neutral, so that there are six treatments instead of two. In practice, the resources may not permit this elaboration, and the issues that must be faced are how best to use the resources available, and whether the experiment, as it can be done, is worth doing. Here again, the deciding factor is usually the use to which the results are to be put.

4. Discussion may arise as to the need for a "control," this term being used rather loosely for a treatment in which we are not particularly interested, but which may be needed in order to reveal, by comparison, whether other treatments are effective. Suppose that we wish to com-

pare the effectiveness of three treatments that are qualitatively similar, for example, three nitrogenous fertilizers all of which supply the same amount of nitrogen. The control would be a "no-nitrogen" treatment. Three cases may be distinguished. (i) The effectiveness of this type of treatment has been consistently demonstrated previously, and it remains to discover which of the three qualities is best. There is no need for a control. (ii) The type of treatment is in general effective, but occasionally the conditions of the test are such that it is not. For example, nitrogenous fertilizers may fail to produce responses in fields where the soil fertility is very high. In this case it will be well to add a control, which serves primarily to describe the conditions under which the fertilizers were tested. (iii) It may not be known whether the type of treatment is effective. The control should of course be included, and there may be a case for giving it more replication than the other treatments receive. An experiment of this type is analyzed in chapter 3, where the treatments were 4 types of soil fumigant, and the control (no fumigation) was replicated 16 times as against 4 for the other treatments. The effect is to obtain a more accurate estimate of the *average* response to fumigants, at the expense of some loss in accuracy in the comparisons among the fumigants.

An interesting example of the illumination that is sometimes produced by the inclusion of a control has been reported by Jellinek (1.3). A headache remedy contained 3 ingredients, *a*, *b*, and *c*. In order to test whether ingredients *b* and *c* were necessary, the complete mixture (*abc*) was compared with (*ac*) and (*ab*). There were 199 subjects, each of whom was treated with each drug for a two-week period, the appropriate drug being given whenever the subject complained of a headache. Success was measured by the ratio of the number of headaches relieved to the number tested in the two-week period. The mean success rates were 0.84 for (*abc*), 0.80 for (*ac*), and 0.80 for (*ab*); these figures show very little overall difference in effectiveness.

The experiment also contained a "control"—an inert mixture which looked the same as the others but had no active ingredients. This was tested under the same conditions as the 3 drugs. No less than 120 subjects, or about 60%, reported at least some of their headaches relieved by the control, a result that is itself of some interest. Further, the control enables us to isolate the group of 79 subjects whose headaches were not relieved by an inert mixture. For this group, the mean success rates were 0.88 for (*abc*), 0.67 for (*ac*), and 0.77 for (*ab*), and could be shown to be significantly different. As the author comments: "Banal as it may sound, discrimination among remedies for pain can be made only by subjects who have a pain on which the analgesic action can be tested."

If a control is required, it must be an integral part of the experiment so that results for the control are directly comparable with those for the other treatments. This point tends to be overlooked in experiments with human subjects when it is difficult or troublesome to assemble the desired number of subjects. For example, if a new drug is to be tested in some ward of a hospital, the recovery rate in the ward before the drug was introduced is not a satisfactory control, nor is the recovery rate in a different ward where patients happen to be receiving the standard drug. An observed difference between the effects of the new and the standard drug might be due to differences in the severity of the disease or in the type of patient or in other aspects of the medical care in the two time-periods or the two wards. It is necessary to regard the new and the standard drug as two experimental treatments on an equal footing, and to use randomization, or one of the methods of restricted randomization, in assigning the two drugs to patients.

1.24 Further Details. Other features of the experiment that should be included in the draft of the proposals are the number of replications, the types of experimental material to be used, and the measurements that are to be made. Since these primarily affect the accuracy of the experiment, discussion of them is deferred to chapter 2, which is devoted to the general question of accuracy.

Finally, the draft should describe in some detail the proposed method for drawing conclusions from the results. This is the section that is most frequently omitted, though it is a valuable one. It may include a sketch of the analysis of variance, an indication of the tabular form in which results will be shown, and some account of the tests of significance to be made and the treatment differences that are to be estimated. In this process we verify which treatment comparisons are relevant to each of the stated objectives of the experiment. Attention is drawn to deficiencies in the set of treatments and to treatments that supply no information essential to the purpose of the experiment.

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CHAPTER 2

METHODS FOR INCREASING THE ACCURACY OF EXPERIMENTS

2.1 Introduction

The results of experiments are affected not only by the action of the treatments, but also by extraneous variations which tend to mask the effects of the treatments. The term experimental errors is often applied to these variations, where the word *errors* is not synonymous with "mistakes," but includes all types of extraneous variation. Two main sources of experimental errors may be distinguished. The first is inherent variability in the experimental material to which the treatments are applied. We shall use the term *experimental unit* to denote the group of material to which a treatment is applied in a single trial of the experiment. The unit may be a plot of land, a patient in a hospital, or a lump of dough, or it may be a group of pigs in a pen, or a batch of seed. It is characteristic of such units that they produce different results even when subjected to the same treatment: these differences, whether large or small, contribute to the experimental errors. The second source of variability is lack of uniformity in the physical conduct of the experiment, or in other words, failure to standardize the experimental technique.

Neither the presence of experimental errors nor their causes need concern the investigator, provided that his results are sufficiently accurate to permit definite conclusions to be reached. In many fields of research, however, with the time and labor that can be given to an experiment, the results are so greatly influenced by experimental errors that only treatment differences that are large can be detected, and even these may be subject to considerable uncertainty. As a consequence, the investigation of methods for increasing the accuracy of experiments has played a prominent part in experimental research in recent years.

The methods may be classified broadly into three types. The first is to increase the size of the experiment, either through the provision of more replicates or by the inclusion of additional treatments. The second is to refine the experimental technique. Thirdly, we may handle the

experimental material so that the effects of variability are reduced. This may be done by careful selection of the material, by taking additional measurements that provide information about the material, or finally by skillful grouping of the experimental units in such a way that the units to which one treatment is applied are closely comparable with those to which another treatment is applied. These various methods are discussed in more detail in succeeding sections.

At this point it is advisable to discuss briefly the concepts *accuracy* and *precision*. Although these words are regarded as roughly synonymous in most dictionaries, they have sometimes been assigned technical meanings, particularly in the natural sciences and engineering. The difference between the concepts may be illustrated by an experiment in which children are weighed on a machine that has a bias of 1 lb. That is, if the true weight of a child is 46 lb., repeated weighings of the child give readings which vary around 47 lb. The *accuracy* of a measurement of the weight of the child signifies the closeness with which the measurement approaches the true weight, 46 lb. The term *precision*, on the other hand, is concerned merely with the repeatability of the measurements. Thus the precision of a measurement denotes the closeness with which the measurement approaches the average, 47 lb., of a long series of measurements made under similar conditions. It is clear that, if the bias is large, a measurement may be of high precision but of low accuracy.*

Most of the devices mentioned in this chapter, including replication, additional measurements, and skillful grouping of the experimental material, operate on the precision of the experiment. In so far as the method of measurement remains biased, these procedures do not affect this bias. A refinement in technique may reduce it. Other things being equal, an increase in precision is accompanied by an increase in accuracy, though if a large bias is present a substantial increase in precision may result in a trivial increase in accuracy.

For simplicity in presentation, we shall sometimes speak as if no bias is present, that is, as if precision and accuracy are identical. Thus the phrase "true difference between two treatments" will mean the true difference as recorded by the measuring device that was actually used. This will be identical with the natural concept of a "true difference" only if the measurements are not biased in favor of one of the treatments.

* For a discussion of the difficult problem of measuring precision and accuracy, see Shewhart (2.8).

2.2 Number of Replications

Whatever the source of the experimental errors, replication of the experiment steadily decreases the error associated with the difference between the average results for two treatments, provided that precautions (such as randomization) have been taken to ensure that one treatment is no more likely to be favored in any replicate than another, so that the errors affecting any treatment tend to cancel out as the number of replications is increased. The rate at which the experimental error is reduced is predictable from statistical theory. The basic quantity used to measure experimental errors is the error variance per experimental unit, which is defined as the expected value of the square of the error that affects the observation for a single experimental unit. The square root of this quantity is called the *standard error per unit*. If σ^2 is the error variance per unit and there are r replications, the error variance of the difference between the means for two treatments is $2\sigma^2/r$, and the corresponding standard error is $\sqrt{2\sigma^2/r}$. This result is valid unless increased replication necessitates the use of less homogeneous experimental material or of a less careful technique.

In succeeding sections some advice is given on the choice of the number of replications for an experiment. To present realistic advice is not easy. Often the size of an experiment is limited by lack of resources or by the conflicting claims of other experiments. However, even if it is realized that an experiment must fall short of the precision desired, it is a good practice to try to estimate the degree of precision that will be attained and to present this information as part of the proposal for the experiment.

2.21 Number of Replications for Tests of Significance. Consider the difference between the effects of a pair of treatments, which might be a standard procedure and some new procedure which it is hoped will prove better than the standard. The precision desired in an experiment may be specified in either of two ways. We may specify the size of the true difference which the experiment is to *detect* by means of a test of significance, or we may specify how closely we wish to *estimate* the true difference, by stating the width of the confidence interval that we would like to have for the true difference.

Although the specification used is to some extent a matter of individual taste, the approach by means of a test of significance is helpful mainly in the initial stages of a line of research. The reasoning in this approach is roughly as follows. If it can be established that the new procedure is superior to the standard procedure by at least some stated amount, say

20% of their mean, then we will have discovered a useful result. On the other hand, if the first experiment shows no significant difference between the two treatments, we will be discouraged from further investigation of the new treatment. Consequently, the first experiment must be large enough to ensure that if the true difference is 20% or more, it is highly probable that we will obtain a significant difference. Note that in this approach we do not insist that the first experiment give a precise estimate of the true difference, but merely that it give a significant result at the chosen level of significance.

The probability of obtaining a significant result depends on the true standard error σ per unit, the number of replications r , and the number of degrees of freedom n that the experiment provides for estimating the error variance. The exact calculation of the probability is rather complicated; tables have been given by Neyman *et al.* (2.1) and others. The argument used in the following example, though logically faulty, leads to an approximation that is good enough for most practical purposes.

Example. The true standard error per unit is 12% of the mean of the experiment, and there are 8 replicates and 28 d.f. for the estimate of error variance. What is the probability of obtaining a significant result when the true difference is 20%?

Let d be the observed difference and δ the true difference between the mean of the new treatment and that of the standard treatment. The true standard error of d is

$$\sqrt{\frac{2\sigma^2}{r}} = \sqrt{\frac{2(12)^2}{8}}; \text{ i.e., } = 6\%$$

In practice we do not know σ , and in its place we use the estimated standard error per unit s , obtained from the analysis of variance of the results. We will assume that s happens to be equal to σ .

The test of significance of d is made by means of a Student t -test, with 28 d.f. To find the value that d must attain in order to be significant, we multiply the standard error of d , 6%, by the significant value of t . Suppose that the test is made at the 5% significance level, and that it is a one-tailed test.* The 10% value of t in the standard tables, for

* A one-tailed test is used when it is known that the new treatment must be at least as good as the standard procedure. The difference d is declared significant only if d is sufficiently large and positive; if d is negative we ascribe the result to experimental errors, since we do not entertain the possibility that the new treatment could be inferior. A two-tailed test is made when we do not know which treatment is better; large values of d , whether positive or negative, are declared significant. All standard tables of the significance levels of t are computed for two-tailed tests. For a one-tailed test we read the value of t corresponding to twice the significance probability.

28 d.f. is 1.701, so that if d is to be significant we must have

$$d \geq \sqrt{\frac{2}{r}} st_1 = (6)(1.701) = 10.206$$

The problem is to find the probability that this happens.

The quantity

$$\frac{d - \delta}{\sqrt{\frac{2}{r}} s} = \frac{d - 20}{6} = t_2$$

follows the t -distribution with 28 d.f. But it is evident that

$$d \geq 10.206 \quad \text{if and only if} \quad t_2 = \frac{d - 20}{6} \geq \frac{10.206 - 20}{6} = -1.632$$

Hence the probability which we seek is the probability that a value of t , with 28 d.f., should exceed -1.632 . In the table of t , the probability that a value lies outside the limits ± 1.632 is about .11. The probability wanted is therefore $[1 - (\frac{1}{2})(0.11)]$, or about .945. It is highly likely that a significant difference would be found in this experiment.

The flaw in the argument is the assumption that the value of s can be assumed equal to σ , whereas s is actually a random variable. If the argument is carried through algebraically, it gives the following rule for calculating the approximate probability P of obtaining a significant result.

Step 1. Find t_2 from the relation

$$\delta = \sqrt{\frac{2}{r}} \sigma(t_1 + t_2) \quad (2.1)$$

where δ = true difference that it is desired to detect.

σ = true standard error per unit.

r = number of replications.

t_1 = significant value of t in the test of significance (with care to distinguish between one-tailed and two-tailed tests).

Step 2.

$$P = 1 - (\frac{1}{2})p_2 \quad (2.2)$$

where p_2 is the probability corresponding to t_2 in the ordinary t -table. The degrees of freedom in t_1 and t_2 are those available for the estimate of error variance.

TABLE 2.1 NUMBER OF REPLICATIONS REQUIRED FOR A GIVEN PROBABILITY OF OBTAINING A SIGNIFICANT RESULT

Upper figure: Test of significance at 5% level, probability 80%

Middle figure: Test of significance at 5% level, probability 90%

Lower figure: Test of significance at 1% level, probability 95%

One-tailed tests

True difference (δ) as percent of the mean	True standard error per unit (σ) as percent of the mean															
	2	3	4	5	6	7	8	9	10	11	12	14	16	18	20	
5	3	6	9	13	19	25	33	41	50							
	4	7	12	18	26	35	45									
	7	13	22	33	47											
10	2	2	3	4	6	7	9	11	13	16	19	25	33	41	50	
	2	3	4	5	8	9	12	15	18	22	26	35	45			
	3	4	7	9	13	17	22	27	33	40	47					
15	2	2	2	3	3	4	5	6	7	8	9	12	15	19	23	
	2	2	3	3	4	5	6	7	9	10	12	16	21	26	31	
	2	3	4	5	7	8	10	13	15	18	22	29	37	47		
20	2	2	2	2	2	3	3	4	4	5	6	7	9	11	13	
	2	2	2	2	3	3	4	5	5	6	7	9	12	15	18	
	2	3	3	4	4	5	7	8	9	11	13	17	22	27	33	
25	2	2	2	2	2	2	3	3	3	4	4	5	6	7	9	
	2	2	2	2	2	3	3	3	4	5	5	7	8	10	12	
	2	2	3	3	3	4	5	6	7	8	9	11	14	18	22	
30	2	2	2	2	2	2	2	2	3	3	3	4	5	6	7	
	2	2	2	2	2	2	3	3	3	4	4	5	6	7	9	
	2	2	2	3	3	3	4	4	5	6	7	8	10	13	15	

The rule may be inverted in order to give the number of replications r required for a given probability P of obtaining a significant result. In this form we have

$$r \geq 2 \left(\frac{\sigma}{\delta} \right)^2 (t_1 + t_2)^2 \quad (2.3)$$

where δ = true difference that it is desired to detect.

σ = true standard error per unit.

t_1 = significant value of t in the test of significance.

t_2 = value of t in the ordinary table corresponding to $2(1 - P)$.

TABLE 2.1 (Continued)

Two-tailed tests

True difference (δ) as percent of the mean	True standard error per unit (σ) as percent of the mean														
	2	3	4	5	6	7	8	9	10	11	12	14	16	18	20
5	4	7	11	17	24	32	41								
	5	9	15	22	31	42									
	7	14	24	38											
10	2	3	4	5	7	9	11	14	17	20	24	32	41		
	2	3	5	7	9	12	15	18	22	27	31	42			
	3	5	7	11	14	19	24	30	37	45					
15	2	2	3	3	4	5	6	7	8	10	11	15	19	24	29
	2	2	3	4	5	6	7	9	11	13	15	19	25	31	39
	3	3	4	6	7	9	12	14	17	21	24	33	42		
20	2	2	2	3	3	3	4	5	5	6	7	9	11	14	17
	2	2	2	3	3	4	5	6	7	8	9	12	15	18	22
	2	3	3	4	5	6	7	9	11	12	14	19	24	30	37
25	2	2	2	2	2	3	3	3	4	4	5	6	7	9	11
	2	2	2	2	3	3	4	4	5	5	6	8	10	12	15
	2	2	3	3	4	5	5	6	7	9	10	13	16	20	24
30	2	2	2	2	2	2	3	3	3	4	4	5	6	7	8
	2	2	2	2	2	3	3	3	4	4	5	6	7	8	11
	2	2	3	3	3	4	4	5	6	6	7	9	12	14	17

Notes. In constructing the table, it was assumed that the number of degrees of freedom for error is $3(r-1)$; this would apply in a randomized blocks experiment with 4 treatments.

No entries are given where more than 50 replications are required.

The use of this equation is slightly tricky, because the number of degrees of freedom in t_1 and t_2 depends on r . Trial and error may be used until the smallest r satisfying (2.3) is found.

Example. In the previous example, how many replications are required in order to have a chance of 4 in 5 that a significant result will be obtained? Here we have $\delta = 20$, $\sigma = 12$, and we will suppose that the experiment has 4 treatments in randomized blocks, which gives $n = 3(r-1)$ degrees of freedom for the estimate of error. In the t -table, t_1 is the value for probability .10, while t_2 is the value for probability $2(1 - .80)$ or .40.

In order to start the calculation some guess must be made about the value of n . It does not matter greatly if the guess is inaccurate; we will try $n = 30$. For this,

$$t_1 = 1.697; \quad t_2 = .854; \quad \text{so that} \quad (t_1 + t_2) = 2.551$$

Hence

$$r \geq 2(12\frac{1}{20})^2(2.551)^2 = 4.7$$

Consequently our second trial value is $r = 5$. Since this provides only 12 instead of 30 d.f., we must repeat the calculation to verify whether 5 replications are sufficient. For 12 d.f. $(t_1 + t_2)$ is 2.655. Thus

$$2\left(\frac{\sigma}{\delta}\right)^2(t_1 + t_2)^2 = 2\left(\frac{12}{20}\right)^2(2.655)^2 = 5.08$$

Since this is greater than 5, we conclude that 5 replications are not quite enough, and that 6 is the smallest number of replications that will satisfy the conditions. We can be sure without further computation that 6 will satisfy the conditions, because the right-hand side of the inequality always decreases when we increase r .

Table 2.1, computed by this method, shows the minimum number of replications for a range of values of δ and σ , and is intended to be used in the planning of experiments. Usually it is most convenient to think of δ and σ expressed in percentages of the mean per observation for the experiment, and the range in the table has been constructed with this in mind. Since the value of r depends only on the ratio σ/δ , it does not matter what units are used for δ and σ , provided that they are the same for both. There are three entries for any given pair of values of δ and σ . These show, in descending order, the replications required for a test of significance at the 5% level and a probability .80 of getting a significant result, for a 5% level and probability .90, and for a 1% level and probability .95. The top half of the table applies to one-tailed tests, the lower half to two-tailed tests.

Construction of the table requires some assumption about the size of the experiment. In the case presented the experiment has four treatments in randomized blocks, so that $n = 3(r - 1)$. The values in the table change little if the experiment happens to provide more than this. If the experiment has only two or three treatments it is advisable to check the value of r by means of the formula.

In the practical use of the table it is necessary to estimate the value of the true standard error per unit. Often this can be done fairly well from

the results of previous experiments with the same kind of material. It should be realized that the probabilities to which the table applies will be correct only if the value of σ is estimated correctly. If σ is underestimated, the probability of obtaining a significant result will of course be smaller than the presumed value. For cases where it is highly important that the probability be correct, more exact methods are discussed in section 2.23.

As an illustration of the reading of the table, consider the case in the last example, where $\delta = 20$, $\sigma = 12$, $P = .80$, and the test is a one-tailed test at the 5% level. For this we read the top entry in the top half of the table for $\delta = 20$, $\sigma = 12$, and find that 6 replications are necessary. From the middle figure we see that an increase to 7 replications increases the probability to .90. If the test is to be two-tailed, we read at the corresponding place in the lower half of the table, and find that 7 replications are required for a probability .80 and 9 for a probability .90.

The table brings out the value of any reduction in the standard error per unit. In fact, we cannot have a high probability of detecting a 5% difference with any reasonable number of replications unless the standard error per unit is 4% or under. Differences of 20% or over can be detected in most cases without excessive replication. No entries have been inserted in the table for situations where more than 50 replications would be required, since experiments of this size are uncommon.

2.22 Number of Replications for Prescribed Limits of Error. The confidence limits for δ , the true difference between the effects of two treatments, are

$$\delta = d \pm \sqrt{\frac{2}{r}} st \quad (2.4)$$

where t is the value in the t -table corresponding to the confidence probability chosen and to the number of degrees of freedom n in the estimate of error. The quantity $\sqrt{2st}/\sqrt{r}$ may be called the *limit of error*, since it measures the maximum distance between d and δ for the confidence probability chosen. Since the limit of error depends on s , it is not known until the experiment is completed. If s is replaced by its average or expected value, we obtain the *expected limit of error*

$$L = \sqrt{\frac{2}{r}} f \sigma t \quad (2.5)$$

The factor f enters because the average value of s is slightly less than σ ; to a close approximation, f equals $\left(1 - \frac{1}{4n}\right)$.

From relation (2.5) table 2.2 was constructed to show the numbers of replications required for various values of L and for confidence probabil-

TABLE 2.2 NUMBER OF REPLICATIONS REQUIRED FOR GIVEN LIMITS OF ERROR IN THE DIFFERENCE BETWEEN TWO TREATMENTS

Upper figure: Confidence probability .80

Middle figure: Confidence probability .90

Lower figure: Confidence probability .95

Limit of error ($\pm L$) as percent of the mean	True standard error per unit (σ) as percent of the mean															
	2	3	4	5	6	7	8	9	10	11	12	14	16	18	20	
3	3	4	7	10	14	19	24	31	37	45						
	4	6	11	16	23	31	40	50								
	5	9	15	23	32	43										
4	2	3	4	6	8	11	14	18	21	26	31	41				
	3	4	6	9	13	18	23	29	35	42	50					
	3	6	9	13	19	25	32	40	50							
5	2	2	3	4	6	7	9	12	14	17	20	27	35	44		
	2	3	5	6	9	12	15	19	23	27	32		44			
	3	4	6	9	12	16	21	26	32	39	46					
6	2	2	3	3	4	5	7	8	10	12	14	19	24	31	38	
	2	3	4	5	6	8	11	13	16	19	23	31	40	50		
	2	3	5	7	9	12	15	19	23	27	32		43			
8	2	2	2	2	3	3	4	5	6	7	8	11	14	18	21	
	2	2	3	3	4	5	6	8	9	11	13	18	23	29	35	
	2	3	3	4	6	7	9	11	13	16	19	25	32	40	49	
10	2	2	2	2	2	3	3	4	4	5	6	7	9	12	14	
	2	2	2	3	3	4	5	5	6	8	9	12	15	19	23	
	2	2	3	3	4	5	6	7	9	11	12	16	21	26	32	

ities of .80, .90, and .95. As in table 2.1, the experiment was assumed to have $3(r-1)$ degrees of freedom in the estimate of error; no entries were inserted where more than 50 replications are necessary, and the values of L and σ are expressed as percentages of the mean for the experiment.

This table may be used in the planning of an experiment where the object is to estimate the difference in the effects of two treatments. Suppose that the true standard error per unit is expected to be about 6%, and that preliminary experiments have indicated that the true difference between the effects of the two treatments is around 25%. In this case we would probably not be satisfied with a value of L any higher than 5%; if L is 5% and if the observed difference is found to be 22%, we are confident that the true difference lies between $(22 \pm 5)\%$, that is, between 17 and 27%. From the entry for $\sigma = 6$, $L = 5$, we find that 6 replications suffice for a confidence probability of 80%, 9 for 90%, and 12 for 95%.

It is important to reiterate that this table does not *guarantee* that the limits of error will be of the desired size, even if the value of σ is forecasted correctly. The actual limits will depend on s and may be either larger or smaller than L . They will be about the desired size *on the average* if σ is guessed correctly. Further, the table makes it clear that small treatment differences of the order of 5% cannot be accurately estimated in a single experiment, at least with the values of σ and r that normally prevail in many lines of research. For an accurate estimate of a 5% true difference, we would probably not want L any larger than 1%. This value was not included in the table since it demands great numbers of replications.

2.23 The Case Where Additional Assurance Is Desired. We have seen that tables 2.1 and 2.2 do not ensure that the desired standards of accuracy will always be attained. In table 2.1 the probability of obtaining a significant result is as stated only if the value of σ is correctly guessed. In table 2.2, if σ is correctly guessed, the limits of error will equal the desired L on the average, but for a single experiment may be either higher or lower. In certain circumstances, methods are available which provide a more definite assurance.

If an initial estimate s_1^2 of σ^2 , based on n_1 degrees of freedom, has been obtained from preliminary data, Harris, Horvitz, and Mood (2.3) present tables that give the number of replications required so as to be certain that the probability of obtaining a significant result will have the desired value. No guesswork about σ is involved, though if n_1 is small the number of replications required tends to be large so as to allow for the possibility that s_1^2 is a very poor estimate of σ^2 . With the same preliminary data a value of r may be found such that the probability is .05 that the limits of error exceed L . If in default of an estimate s_1^2 we can give a range of values within which σ^2 is likely to lie (e.g., from pre-

vious experiments), the authors suggest a simple method for using their tables.

An alternative approach, due to Stein (2.4), is applicable when the experiment can be interrupted after a given number of replications. This approach ensures that the limits of error will not exceed L . The procedure is as follows. We first complete a number of replications somewhat less than the number which we expect will be necessary. At this stage we compute the limits of error. If these are already less than L , the object is achieved. If they are greater than L , Stein gives a simple rule for computing the additional number of replications needed to give limits that are less than L .

Naturally, these methods require more replications than tables 2.1 and 2.2. In fact, if used repeatedly, they give a standard of accuracy which on the average is greater than we desire, in order to ensure that the standard is seldom if ever less than what is wanted. For this reason, tables 2.1 and 2.2 appear to be suitable for ordinary experimentation; the methods described here should be used when it is very important to reach a specified standard of accuracy.

2.3 Other Methods for Increasing Accuracy

2.31 The Measurement of Relative Efficiency. We have seen that the error variance of the difference between the means for two treatments is $2\sigma^2/r$, where r is the number of replications. This result provides a simple means for comparing, in concrete terms, the relative precision of two experiments. Suppose that one design gives a true error variance of 1.0 per experimental unit, while a second gives 0.5. The two experiments would give equally precise comparisons among the treatment means if the amount of replication in the first were twice that in the second. From a knowledge of the error variances per unit, comparisons are thus easily made in terms of the relative amounts of replication required to attain the same degree of precision. The inverse ratio of the variances per unit is sometimes called the *relative efficiency* of the two designs; in the example above, the relative efficiency of the second experiment to the first is 2.0 or 200%. Similarly we might speak of the *relative gain in efficiency* as 1.0 or 100% in this case. These terms occur frequently in the literature on experimental designs.

One further factor must be taken into account. For the same numbers of treatments and replicates, the number of degrees of freedom in the estimate of error changes with the design. This number enters into many of the uses to which the experimental results are put. When the number of degrees of freedom becomes smaller, the limits of error for a true

difference are increased and the probability of obtaining a significant result is decreased; in other words, the sensitivity of the experiment is decreased.

Thus the numbers of degrees of freedom for error are relevant to the comparison of two designs. A change in design which decreases the error degrees of freedom as well as the error variance may not be advantageous. This leads to the question: by how much must the experimental error variance be reduced so as to balance a given reduction in the error degrees of freedom? The question has been discussed by Neyman *et al.* (2.1), Fisher (2.5), and Walsh (2.6). The answer depends on the use to which the results are put. From one point of view, two experiments may be regarded as equal in sensitivity when the limits of error for a given confidence probability are equal. Alternatively, equal sensitivity may imply an equal probability of detecting as significant a given real difference between the effects of two treatments. Other definitions could be set up.

Table 2.3 illustrates the situation. An experiment with a true error variance per unit equal to 1 (i.e., with an infinite number of degrees of freedom) was chosen as the standard. The table shows, for a range of

TABLE 2.3 EFFECT OF NUMBER OF DEGREES OF FREEDOM IN ERROR ON THE SENSITIVITY OF THE EXPERIMENT. ERROR VARIANCES PER UNIT WHICH GIVE EQUAL SENSITIVITY ACCORDING TO FOUR DEFINITIONS

Definition	Number of error degrees of freedom (n)											
	1	2	3	4	5	6	8	10	15	20	30	∞
1	0.04	0.26	0.45	0.56	0.64	0.70	0.77	0.81	0.87	0.90	0.94	1.00
2	0.27	0.59	0.72	0.79	0.83	0.86	0.90	0.92	0.94	0.96	0.97	1.00
3	0.22	0.47	0.61	0.69	0.75	0.79	0.84	0.86	0.91	0.93	0.95	1.00
4	0.10	0.40	0.57	0.67	0.73	0.78	0.83	0.87	0.91	0.93	0.96	1.00

values of the error degrees of freedom (n), the true error variances per unit which are required to equal the standard in sensitivity according to four definitions of this term. In case 1 the average limits of error for a confidence probability .95 are made equal. In case 2 the same criterion is used for a confidence probability .80. In case 3 all experiments have the same probability .25 of detecting as significant a constant difference between two treatments. Case 4 shows the same comparison for

a larger constant difference where the probability is .75. The tests of significance are one-tailed t -tests at the 5% level.

For example the figure 0.70 for $n = 6$ (definition 1) means that with a true error variance per unit of 0.70 and 6 d.f. in the estimate of error, the limits of error for a confidence probability .95 are the same on the average as when σ^2 is 1 and n is infinite. A 30% decrease in error variance is needed to compensate for the lack of degrees of freedom.

The wide divergence between the results for definition 1 and definition 2 is noteworthy since the only change in criterion is that from a confidence probability .95 to one of .80. Definitions 3 and 4 exhibit much closer mutual agreement and give results intermediate between those of definitions 1 and 2. In practical applications the discrepancies among the four methods will be smaller than table 2.3 suggests, because comparable designs for the same experiment seldom differ greatly in the degrees of freedom available for error.

Fisher's approach (2.5) is somewhat different. In effect, he calculates the "amount of information" which the estimated difference d between two treatment means supplies about the true difference δ , the calculation being made from the t -distribution of the quantity

$$\frac{\sqrt{r}(d - \delta)}{\sqrt{2s}}$$

He finds the information to be $(n + 1)/(n + 3)s^2$, whereas if σ were known exactly the information would be $1/\sigma^2$. The estimated variances which give equal amounts of information are shown in table 2.4.

TABLE 2.4 ESTIMATED VARIANCES WHICH PROVIDE EQUAL AMOUNTS OF INFORMATION

Number of error degrees of freedom (n)															
1	2	3	4	5	6	7	8	9	10	11	12	15	20	30	∞
0.500	0.600	0.667	0.714	0.750	0.778	0.800	0.818	0.833	0.846	0.857	0.867	0.889	0.913	0.939	1

This table agrees closely with the results for cases 3 and 4, except when n is below 5. For general purposes it is suggested that this table be used to take account of the differences in degrees of freedom for error in two designs that are being compared. Suppose that a design with $n = 6$ is compared with one in which $n = 12$. The former gives more information only if the error variance s_1^2 is less than $(0.778/0.867)s_2^2$.

Thus the relative efficiency of the *first* design to the *second* is estimated as $0.778s_2^2/0.867s_1^2$, or more generally as

$$\frac{(n_1 + 1)(n_2 + 3)s_2^2}{(n_2 + 1)(n_1 + 3)s_1^2}$$

The adjustment is of importance only if n_1 and n_2 are small.

This expression applies to the simple case in which the two experiments have the same set of treatments. The experiments may, however, have *different* sets of treatments, although each provides an estimate of some specific treatment effect in which we are interested. Alternatively, a different method of estimation of the treatment effect may have been employed in the two experiments. In such cases we may wish to assess the relative accuracy with which this treatment effect is estimated in the two experiments. The expression given above still applies, except that s_1^2 and s_2^2 become the estimated variances of the treatment effect as found in the two experiments, while n_1 and n_2 are as before the numbers of degrees of freedom in s_1^2 and s_2^2 respectively.

In succeeding sections a number of other devices which affect the accuracy of an experiment are presented briefly.

2.32 Selection of Treatments. In certain cases the selection of the treatments has a substantial effect on the precision of an experiment. As Fisher has stressed, striking gains in precision may be achieved by testing different types of treatment in the same experiment, instead of conducting a separate experiment for each type. For instance, one type of treatment or *factor* might be the depth of ploughing in the preparation of a field for a crop, while another might be the addition of a nitrogenous fertilizer. We might conduct one experiment in which deep (*D*) and shallow (*S*) ploughing are compared, and another in which no fertilizing (*O*) is compared with the addition of a specified amount of fertilizer (*N*). In a *factorial* experiment both factors would be tested simultaneously by means of the 4 "treatments"—(*DO*), (*SO*), (*DN*), and (*SN*), where the symbol (*DO*) implies deep ploughing with no addition of nitrogen, etc. The average response to nitrogen is assessed by comparing the last two treatments with the first two. This comparison gives as precise an estimate of the average response to nitrogen as if the whole experiment had been devoted to that factor alone. The same property holds for the estimation of the average difference in effect between deep and shallow ploughing. Thus the experiment, as it were, does double duty. The use of factorial experimentation is discussed more fully in chapter 5.

There are many specialized problems where the choice of the proper

amounts of some ingredient is important. As a very simple example, suppose that the response to increasing amounts of the ingredient is known to be linear and that the purpose is to determine the slope of the line. The most accurate experiment contains only two amounts of ingredient. These should be placed at the ends of the range within which the response is linear and experimentation is feasible. With three amounts of ingredient, two at the ends and one in the middle, the variance of the estimated slope, for the same total size of experiment, is $1\frac{1}{2}$ times as large, and if more amounts are used it becomes larger still. A more complex example occurs in the estimation of the amount of a virus in a solution by preparing a series of dilutions of the solution and counting the numbers of lesions produced on leaves which are rubbed with the dilutions (2.7). The accuracy of the estimate is known to depend both on the dilution ratio that is chosen and on the number of dilutions that are used.

2.33 Refinements of Technique. Since *technique* is the responsibility of the experimenter, its importance need not be elaborated. The principal objectives of a good technique are as follows.

i. To secure uniformity in the application of the treatments. In pig-feeding experiments, for instance, a uniform amount of food cannot be supplied to each animal without the provision of individual feeding boxes. In the testing of insect sprays, delicate apparatus is required in order to subject each batch of insects to a desired dose of spray.

ii. To exercise sufficient control over external influences so that every treatment produces its effects under comparable and desired conditions. It is difficult to generalize about the degree of control needed; a balance must be struck between the cost incurred and the gain in precision obtained. The artificial production of diseases for experiments on resistance to infection exemplifies a case where experimentation cannot proceed rapidly without such control over external conditions.

iii. To devise suitable unbiased measures of the effects of the treatments. Often the appropriate measurements are readily apparent; sometimes the development of a satisfactory method of measurement requires years of research, as in the estimation of certain of the vitamins, in soil analysis, and in sociological investigations.

iv. To prevent gross errors, from which no type of experimentation seems to be entirely free. Adequate supervision and checking of the work of assistants and a scrutiny by the experimenter of the data from every experimental unit will go far towards the discovery and rectification of errors.

Faulty technique may increase the real experimental errors in two ways. It may introduce additional fluctuations of a more or less

random nature. Such fluctuations, if they are substantial, should reveal themselves in the estimate of error as calculated in the analysis of variance (chapter 3). Where his estimated standard errors are consistently higher than those of other workers with similar material, the experimenter is advised to seek the reason, which may lie in differences in technique. In addition, faulty technique may result in measurements that are consistently biased. The estimate of error does not take account of such biases, since it is derived from comparisons of the measurements with one another; in other words, it estimates precision rather than accuracy. The principal safeguards against such biases are care and skill in the construction and handling of measuring devices, plus the intelligent use of randomization.

It is worth while to consider from time to time whether simplifications can be brought into the technique without undue loss of accuracy. Many experiments involve chemical determinations which vary little from unit to unit within the same treatment. For instance, in sugar-beet experiments, the experimental error of the amount of sugar per acre is due mainly to the error in the weight of roots per plot. The amount contributed by the sugar content percent is nearly negligible.

With compound measurements of this type, the labor devoted to the less variable component can be reduced with little loss of accuracy. If the chemical measurement is expensive, a considerable saving in cost may be made. Tests of significance of the chemical composition need not be sacrificed. For example, if the experiment has six replicates, a chemical analysis might be made for each treatment on bulked material from the first three replicates and another on bulked material from the last three replicates. These data provide a valid test of significance based on two "replicates."

2.34 Selection of Experimental Material. The choice of the experimental unit may be of importance. In the planning of field experiments numerous studies have been made of the variability among crop yields on plots of various sizes and shapes under uniform treatment. From these data the best size and shape are selected. The criterion should be to obtain the maximum accuracy for a given expenditure of time and labor.

Frequently, uniform material is prepared specially for experimental purposes, as in the development of inbred lines by laboratories engaged in animal experimentation. Alternatively, the experiment may be confined to a sample chosen for its homogeneity from a large batch of experimental material. If the results of the experiment are to be applied to unselected material, these types of specialization have potential disadvantages. Responses obtained to the treatments in highly selected

material may not be the same as the responses in unselected material.

Selectivity in the material is difficult to avoid where experiments are conducted in the field of economics or sociology. In testing some method of farm management, for instance, the experiment may require the active participation of a group of farmers. The success of such experiments depends greatly on the tact and resourcefulness of the investigator in persuading farmers to cooperate so that the participants are a representative sample of the population about which generalizations are to be drawn.

2.35 Additional Measurements. In the course of an experiment it may be possible to take supplementary measurements which predict at least to some extent the performances of the experimental units. In an experiment to measure the effects of different diets on the weights of children, their weights at the start of the experiment would be a supplementary measurement of this kind, since the increase in weight of a child during the experiment is probably correlated with his initial weight.

By a technique known as the *analysis of covariance*, we may estimate from the data the extent to which the observations were influenced by the variations in these supplementary measurements. The average response to each treatment can then be adjusted so as to remove the experimental error arising from this source. In the school experiment, the *adjusted* responses to treatments represent approximately the responses that would be obtained if all the children had the same initial weight. Thus the effects of variations in the initial weights are largely eliminated from the experimental error, without the necessity of equalizing the initial weights in the planning of the experiment.

Since the analysis is a statistical technique of comparatively recent origin (it was first presented by Fisher about 1932), it may be unfamiliar to many investigators who could use it to obtain a substantial increase in precision with little effort. Its purpose is to remove experimental errors arising from extraneous sources of variation which it is impractical or too costly to control by a more refined technique. Cases are fairly common where the use of covariance has more than doubled the efficiency of the experiment. The extra work involved consists in taking the subsidiary measurements and in applying the adjustments to the treatment means. An example of the computations is given in section 3.8.

2.36 Planned Grouping: Complete Blocks. Finally, we may attempt to minimize the experimental errors through the choice of the experi-

mental plan. The prospects of increased accuracy by this means have been widely explored during the past twenty years. This book is intended as a source of reference to the numerous procedures that have been developed.

The basic idea is simple. Consider an experiment of which a number of separate replications have been conducted. The experimental errors of the results from any replicate can arise only from sources of variation that affect the units within that replicate. Consequently, the error of the difference between two treatment means taken over a number of replicates must also arise solely from variations *within* the individual replications. Variations from one replicate to another do not contribute to the errors. In carrying out an experiment we utilize this simple fact by trying to control sources of variation that affect different units in the same replicate. We need not attempt to reduce differences among the replicates. For example, if the experimental units form a very heterogeneous batch, we attempt to group them so that units in the same replicate are similar; we do not worry if the units in one replicate are not at all similar to those in another. By this device, precise experiments can often be made from what at first sight appears an unpromising batch of material. Similarly, if a uniform experimental technique cannot be maintained throughout the experiment, the important point is to keep the technique uniform within a replication; changes should be made when moving from one replicate to another. This type of design, known in agriculture as *randomized blocks*, is discussed in section 4.2.

The idea is carried a stage further in the latin square (section 4.3). The treatments are arranged diagrammatically so that each appears once in every row and once in every column of a square array. Variations among the groups of experimental units which correspond to the rows and also among those which correspond to the columns are eliminated from the experimental errors. For example, in a comparison of 5 different ways of performing some manual operation, the rows may represent 5 operators, each of whom carries out all the methods in turn. The columns, numbered from left to right, may prescribe the order in which each operator performs the 5 tasks. The plan of a latin square design for this experiment might read as follows.

Order of procedure					
Operator	(1)	(2)	(3)	(4)	(5)
I	D	A	B	E	C
II	A	B	C	D	E
III	B	E	A	C	D
IV	C	D	E	B	A
V	E	C	D	A	B

The letters *A, B, C, D, E* represent the 5 methods. Thus operator I carries out method *D* first, method *A* next, and so on. With this arrangement, the experimental errors are unaffected by differences among the abilities of the men and also by systematic variations introduced by the order in which the methods are performed. If the opportunity presents itself, this type of control may include a third factor by the use of a graeco-latin square (section 4.5).

2.37 Planned Grouping: Incomplete Blocks. In the randomized block design, the precision depends to some extent on the number of treatments. As this number increases, it becomes more difficult to keep the variations between experimental units within each replicate small, and the error variance per unit tends to increase. In experiments with a large number of treatments, numerous attempts have been made to avoid this loss of precision by the formation of groups of experimental units which do not contain all the treatments and therefore can remain small. These groups are called *incomplete blocks*. The groups are so constructed that we can still remove the effect of the differences among groups from the experimental errors, just as such differences are automatically removed in the randomized blocks design. With incomplete blocks, however, this removal usually requires the application of adjustments to the treatment means, so that the statistical analysis of the results becomes more complex.

The best method of constructing the incomplete blocks varies with the nature of the experiment. Sometimes certain comparisons among treatments are of greater interest than others. For instance, in a field experiment containing a number of early-maturing and an equal number of late-maturing varieties of a crop, comparisons between pairs of varieties which mature at about the same date will presumably be more important than comparisons between an early and a late variety. An incomplete block might contain either all the early or all the late varieties, a pair of blocks forming a complete replication. This arrangement, of which several variations are possible, is known as the *split-plot* design (section 7.1).

A similar situation occurs in factorial experiments. Certain comparisons, e.g., the average effects produced by varying one factor, are nearly always of immediate interest. Others, which represent rather complex relationships among the effects of different factors, may be of minor importance, often because previous experiments have shown that these interrelationships, called *interactions*, exert little or no influence. It has been found that, by a deliberate sacrifice of precision in the estimates of these interactions, the size of the group or block can be re-

duced so as to produce an increase in the precision with which the average effects of the factors are estimated. This principle, known as *confounding*, can also be combined with the use of the latin square (chapters 6 and 8).

If all comparisons between pairs of treatments are potentially of equal importance, a different method is used in forming the blocks. When treatments are arranged in incomplete blocks, two treatments which occur in the same block are more precisely compared than two which are placed in different blocks, at least in so far as the investigator has succeeded in decreasing the variation within blocks. Therefore the goal is to construct a design such that *any pair of treatments* occurs equally often within some block. This type of problem has been investigated previously for its mathematical interest. A solution can be found for any number of treatments and any size of block, but most of the solutions require too many replications for the usual conditions of experimentation. The solutions available to date (called *balanced incomplete block designs* (chapter 11)) are given for all cases in which the number of replications does not exceed ten. Here again the latin square principle can sometimes be used (chapter 13) to eliminate variation amongst two different types of grouping of the experimental units.

For a given number of treatments and a given size of incomplete block, balanced designs allow little choice in the number of replications. Thus, with 64 treatments and blocks of 8 units each, 9 replications are required to balance the design. In order to provide designs for small numbers of replications, a number of additional types have been developed; these are similar to balanced designs except that they lack complete symmetry.

One group, called *lattice* designs (chapter 10), has been used widely in agricultural experiments during the past 5 years. For these designs, the number of treatments must be an exact square, e.g., 25 or 49, while the number of units in the block is the corresponding square root, 5 or 7, respectively. A further group, known as *cubic lattices* (section 10.4), is useful when the number of treatments exceeds 100 yet it is desirable to have a small block. In this case the number of treatments is the cube of the number of units per block, and the number of replications may be any multiple of 3.

2.38 Summary. There are numerous methods available for increasing the accuracy of an experiment. Frequently the same end may be reached in several different ways. Thus, in the experiment which tests the effects of a number of diets on the weights of children, the disturbing effects of variations in their initial weights can be reduced by selecting

for the experiment a number of children of approximately the same initial weight, or by the use of the analysis of covariance, or by a planned grouping in which children within the same replicate are of approximately equal initial weight, or simply by having a large number of replications. The method adopted should be that for which the desired standard of accuracy can be attained with the smallest expenditure of time and effort. There is no special merit in either a complicated experimental plan or a highly refined technique if equally accurate results can be secured with less effort in some other way. A good working rule is to use the simplest experimental design that meets the needs of the occasion. This is not to say that the more complex designs will be used only rarely; in fact, they have already demonstrated their utility in fields of research where no other equally practicable method appears to exist for reaching the same results.

2.4 The Grouping of Experimental Units

2.41 Investigation of Methods for Grouping. In order to take full advantage of the opportunities for increased precision by suitable grouping of the units, the investigator must know the best criteria for grouping. Frequently these are suggested from previous observations on the experimental material or on disturbing factors that become evident in the course of an experiment. Where knowledge is less definite, as in a new line of research, there are two ways in which information may be accumulated.

One is to devote an experiment specifically to this problem. In this case it is usually advisable to subject all units to the same treatment, carrying out a so-called *uniformity trial*. The word *uniformity* refers to uniformity of treatment; the experimental material and technique should be representative of those which are used in actual experiments. The amount of material should be sufficient to enable the experimenter to superimpose on the results a hypothetical trial of the size which he customarily performs. It is worth while to record any auxiliary measurements which might predict the performance of the experimental units. From the results of these trials any proposed grouping of the units can be formed and the amount of variation within and among the groups can be calculated. If it is desired to compare different experimental plans, each may be superimposed on the results of the uniformity trial. By the method indicated in section 2.31, the investigator may estimate the relative numbers of replications that are needed to reach the same degree of precision with the various plans.

Secondly, useful information can sometimes be secured without any

special trial or technique by examination of the results of *actual experiments*. With an experiment which is suitably designed, it is possible to estimate from an analysis of variance of the results what the error variance would have been if any particular grouping had not been used. This estimate, being derived from the same data, is directly comparable with the error variance obtained by the use of the grouping and enables the investigator to evaluate the success of the grouping. Consequently, for experiments in a new line of research, the experimenter may try any method of grouping that *might* be effective, in the knowledge that its appropriateness will be revealed by the experiments themselves. Instructions for performing such comparisons, which are easily made when the analysis of variance has been completed, are given in the notes which accompany each type of design.

2.42 Criteria for Grouping. Many factors have been used as the basis for grouping. In agricultural experimentation it has been repeatedly demonstrated that plots close together tend to be more similar in their yields than plots farther apart. The almost universal practice is to put neighboring plots in the same group, with the groups approximately square in shape whenever practicable. Contiguity frequently forms a good basis for grouping in greenhouse experiments also, where differences may exist along the bench in temperature, sunlight, air currents, or accessibility for watering. Alternatively, in many plant experiments, and more particularly in animal experiments, some characteristic of the plant or animal is much more useful, the physical location of the experimental units throughout the course of the trial having a relatively minor influence on the results. Age, weight, vigor, sex, and genetic constitution are some of the factors most commonly used. The animal itself may constitute the block in cases where the treatments can be applied in succession to the same animal without producing residual effects which obscure the results.

A single grouping, as in the randomized block design, can be used to eliminate simultaneously variation from a number of different sources. For example, suppose that an experiment were planned to investigate whether some treatment applied to rats enabled the animals to withstand subjection to a dose of a poisonous gas. The replication would contain two animals, a treated rat and an untreated rat to be included as a control. Given a sufficient stock of experimental animals, the pairs in any replication might be animals of the same litter (and consequently of the same age), of the same sex, and of approximately the same weight and vigor. If it were convenient to test two animals at the same time, the pair could be put into the gas chamber together, thus eliminating

also the effects of variations in the strength of the dose or in the time for which it was supplied.

The freedom with which the size of block can be varied depends on the type of research. In field experiments the block can be laid down so as to accommodate any number of treatments, though, as previously stated, the standard error per plot may be expected to increase slowly as the size of block increases. When the use of designs with incomplete blocks is contemplated for field experiments, the principal question is to decide for what number of treatments the reduction in block size becomes worth while. On the other hand, the number of leaves on a plant, or of animals of the same litter, usually varies within rather narrow limits. In such cases, where the most homogeneous grouping fixes the size of the block, incomplete block designs may be of special interest.

The gains in efficiency obtainable by skillful construction of the blocks, when measured in terms of increased replication, are often large; examples can be cited in which the effective replication has been increased from two- to tenfold. Frequently, apart from the preliminary care and thought involved, an accurate design uses the same material and involves the same physical operations as the design which it replaced. In some of the more complex designs the computations required to estimate the treatment averages and to perform tests of significance become rather laborious and involved. The experimenter's reaction to these difficulties will depend on his aptitude for numerical computations. At a first trial considerable time may be spent in mastering some of the methods, but the processes become easier with familiarity.

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CHAPTER 3

NOTES ON THE STATISTICAL ANALYSIS OF THE RESULTS

3.1 Introduction

With regard to the statistical analysis for a design, the procedure adopted in this book is to present the computations, with worked numerical examples for the designs likely to be most frequently used. To supply a complete mathematical justification of the analysis for each type of experiment, although in many respects desirable, would have unduly increased the amount of material. The same general technique governs the analysis of all designs, differences arising only in the adaptation of the technique to the particular structure of a design. In the present chapter this technique—the method of least squares—is described and applied to one of the simplest designs. With perseverance the method may be used to verify the computing instructions for other designs in this book. It should be pointed out that the computing methods that are easiest in practice are not necessarily those that flow from a straightforward application of the theory, so that at first sight some of the instructions may appear different from those given by theory.

Some account will also be given of certain statistical techniques, such as the technique for missing values and the analysis of covariance, that are particularly useful in handling the results of experiments, though they are perhaps too specialized to be included as part of a general introductory course in statistics.

3.2 The General Method of Analysis

3.21 The Mathematical Model. The illustrative data come from an experiment conducted in 1935 by the Rothamsted Experimental Station (3.1). The object of the experiment was to measure the effectiveness of 4 soil fumigants in keeping down the numbers of eelworms in the soil. The fumigants were chlorodinitrobenzene (CN), carbon disulphide jelly (CS), and two proprietary preparations, "Cymag" (CM) and "Seekay" (CK). Each fumigant was tested both in a single and a double dose. These comprise 8 distinct treatments. The control (no fumigant) was

the ninth treatment. Normally the replication would have consisted of 9 plots, but in this case 4 control plots were placed in each "replicate," which actually contained 12 plots. The purpose was to supply a fairly accurate standard against which to measure the performance of the fumigants.

The experiment was laid out in 4 separate blocks of 12 plots each, so that there were 4 replications for each dose of each fumigant and 16 replications of the control. Within a block the 12 "treatments" were allotted to individual plots at random. The experiment is of the familiar type known in agriculture as randomized blocks (see chapter 4). The fumigants were ploughed in during spring, after which a crop of oats was sown. After harvest, a sample of 400 grams of soil was taken from each plot and the number of eelworm cysts counted for each sample. The data are those labelled "second count" in table 3.1.

TABLE 3.1 PLAN AND NUMBERS OF EELWORM CYSTS PER 400 GRAMS OF SOIL, FIRST COUNT ABOVE, SECOND BELOW

Block 1 Totals 2587 4383	0	2CK	1CN	1CM	2CM	2CS	2CK	0	Block 2 Totals 964 3075
	269	283	252	212	95	127	80	134	
	466	280	398	386	199	166	142	590	
	1CS	0	0	2CM	1CK	1CN	1CM	0	
Block 3 Totals 1743 4752	138	100	197	263	107	89	41	74	Block 4 Totals 872 2470
	194	219	421	379	236	332	176	137	
	2CS	1CK	0	2CN	0	0	2CN	1CS	
	282	230	216	145	88	25	42	62	
Block 3 Totals 1743 4752	372	256	708	304	356	212	308	221	Block 4 Totals 872 2470
	1CK	0	1CS	2CK	2CK	0	1CK	1CM	
	124	211	194	222	193	209	109	153	
	268	505	433	408	292	352	132	454	
Block 3 Totals 1743 4752	0	2CN	2CS	1CN	0	2CN	2CS	0	Block 4 Totals 872 2470
	102	193	128	42	29	9	17	19	
	363	561	311	222	254	92	28	106	
	2CM	0	1CM	0	1CS	1CN	0	2CM	
Block 3 Totals 1743 4752	162	191	107	67	23	19	44	48	Block 4 Totals 872 2470
	365	563	415	338	80	114	268	298	

In a formal analysis of the results the first step is to set up an equation for every observation. This equation expresses the observation as the sum of four components: (i) a general average about which the

observations are presumed to be fluctuating; (ii) a component representing the effect of the treatment applied; (iii) a component representing certain environmental effects which the design of the experiment enables us to isolate; and (iv) a residual component, representing all other sources that influence the observation, and generally referred to as the "experimental error."

Component (iii) cannot be fully appreciated without some knowledge of the technique employed in the construction of designs. This technique consists in arranging the conduct of the experiment so that, in the subsequent analysis of the results, differences among certain groups of observations can be measured. What is more important, the effects of such differences can be eliminated from the estimates of the treatment effects. If this elimination were not feasible, these differences would contribute to the "experimental error," component (iv), with the consequence that less accurate estimates of the treatment effects would be obtained. In the present example, as will be seen, these differences are differences among the groups of 12 observations that constitute a block. With each design that we describe, the environmental components that are eliminated in this way are pointed out.

We proceed to write down the mathematical model as applied to the example. Instead of numbering the observations from 1 to 48, it is convenient to use a subscript i to denote the treatment applied, j to denote the block in which the observation lies, and k to denote the order within the block. For the plots that receive fumigants, the subscripts i and j are sufficient to define each plot uniquely. The subscript k is needed to distinguish between different control plots within the same block. In this notation the equation for any observation may be written

$$y_{ijk} = \mu + \tau_i + \beta_j + e_{ijk} \quad (3.1)$$

where μ represents the general mean, τ_i the effect of the treatment, β_j that of the block, and e_{ijk} the experimental error.

3.22 Assumptions Made in the Model. There is already implicit in the model one assumption that may not be true in the data obtained from experiments. This assumption is that the treatment effect and the block effect are *additive*. If we take the difference between two observations in the same block, say y_{1jk_1} and y_{2jk_2} , we have, as a deduction from the model,

$$y_{1jk_1} - y_{2jk_2} = \tau_1 - \tau_2 + e_{1jk_1} - e_{2jk_2} \quad (3.2)$$

The block effects have disappeared from the right-hand side; that is, the model implies that the difference between the true effects of two treatments is the same in all blocks. On a block with a high eelworm

infestation a successful fumigant is presumed to reduce the eelworm numbers by the same amount as on a block with a low eelworm infestation. This is unlikely to be so, except as a first approximation. Little systematic study of the accuracy of this approximation in practice has been made. From general experience it appears that the approximation works well in a large number of experiments; nevertheless, cases are not infrequent where the approximation is poor.

In addition, several assumptions are made about the residual effects e_{ijk} . These are taken to be independent from observation to observation, and to be distributed with zero mean and the same variance σ^2 . Further, for tests of significance and the estimation of confidence limits, the e_{ijk} are assumed to follow the normal or Gaussian frequency distribution. In practice, these assumptions are only approximately fulfilled. The consequences of errors in the assumptions will be indicated later (section 3.9).

Apart from changes in detail, the type of model and assumptions presented for this example apply to all experimental arrangements included in this book, except for two types of design for which a more complex model is required. These are the *split-plot* designs (chapter 7), where two or more different error variances are postulated, and *incomplete block designs* (chapter 9), where additional assumptions are made about the nature of the environmental effects β_j .

3.23 Estimation of the Treatment Effects. When the assumptions in the previous section are valid, there is a well-known result in statistical theory that the best estimates of the unknowns μ , τ_i , and β_j are obtained by the *method of least squares*. This method chooses estimates m , t_i , and b_j , respectively, which minimize the sum of squares of the residuals

$$\sum (y_{ijk} - m - t_i - b_j)^2 \quad (3.3)$$

taken over all observations.

In order to find these minimizing values we differentiate the sum of squares with respect to each unknown in turn, and set the derivative equal to zero. Consider a specific unknown, e.g., t_i . If any square contains t_i , its derivative is

$$-2(y_{ijk} - m - t_i - b_j)$$

If the square does not contain t_i , the derivative is zero. Hence, if the derivative of the sum of squares in (3.3) is set equal to zero, the resulting equation may be written

$$\sum (y_{ijk} - m - t_i - b_j) = 0$$

or

$$\sum (m + t_i + b_j) = \sum y_{ijk} \quad (3.4)$$

where the sum is over all observations whose equations contain t_i , or more generally over all observations whose equations contain the parameter to be estimated.

Equations (3.4) are called the equations of estimation, or the *normal* equations. As we have seen, the equation of estimation for any parameter is constructed by equating the total of the observed values y_{ijk} to the total of the expected values $(m + t_i + b_j)$ over all observations whose equation contains the parameter.

We now write down some normal equations for the example. Since m occurs in all 48 observations, its normal equation is

$$48m + 4(t_1 + t_2 + \cdots + t_8) + 16t_9 + 12(b_1 + b_2 + \cdots + b_4) = G \quad (3.5)$$

where t_9 denotes the control treatment, and G the grand total of all 48 observations. At this point a simplification may be introduced. In the model the quantities t_i are needed only to indicate by how much an individual treatment differs from the average of all treatments in the experiment. Accordingly the model is not changed in any essential feature if we assume that the mean of all the t_i is zero. It may also be shown that any convenient linear function of the t_i may be assumed to be zero. The same remarks apply to the block values b_j . From the form of equation (3.5) it appears that convenient assumptions of this kind are

$$t_1 + t_2 + \cdots + t_8 + 4t_9 = 0 \quad (3.6)$$

$$b_1 + b_2 + b_3 + b_4 = 0 \quad (3.7)$$

With these assumptions equation (3.5) reduces to $48m = G$, so that m becomes the mean value of all observations in the experiment.

By summing over the observations that receive a particular treatment t_i we find that its normal equation is

$$4m + 4t_i + b_1 + b_2 + b_3 + b_4 = T_i \quad (i = 1, 2, \dots, 8)$$

and for the control,

$$16m + 16t_9 + 4(b_1 + b_2 + b_3 + b_4) = T_9$$

where the T 's denote observed treatment totals. From relation (3.7), these equations reduce to

$$t_i = \frac{T_i}{4} - m \quad (i = 1, 2, \dots, 8); \quad t_9 = \frac{T_9}{16} - m \quad (3.8)$$

As might be expected, the effect of any treatment is estimated by taking the difference between the mean of the observations which receive that treatment and the general mean.

Let us express these estimates in terms of the original model (3.1). By adding the relevant equations we find

$$\begin{aligned} T_i &= 4\mu + 4\tau_i + (\beta_1 + \beta_2 + \beta_3 + \beta_4) + \sum e_{ijk} \\ T_9 &= 16\mu + 16\tau_9 + 4(\beta_1 + \beta_2 + \beta_3 + \beta_4) + \sum e_{9jk} \end{aligned}$$

Hence, using (3.8),

$$m + t_i = \mu + \tau_i + \frac{1}{4}(\beta_1 + \beta_2 + \beta_3 + \beta_4) + \bar{e}_i$$

where \bar{e}_i is the average of the residual errors on plots that receive treatment i . For the difference between a pair of treatments we deduce

$$t_i - t_u = (\tau_i - \tau_u) + (\bar{e}_i - \bar{e}_u) \quad (3.9)$$

This shows that the estimated difference equals the true difference plus an error of estimate $(\bar{e}_i - \bar{e}_u)$. Note that the error of estimate does not contain any of the environmental (block) effects arising from component (iii). This result exemplifies a valuable property of least squares solutions, namely, that the estimate of the difference between the effects of any two treatments is not influenced by the environmental effects, but only by the residual effects e_{ijk} . It is in this sense that we say that environmental effects are *eliminated* from the estimates of the treatment effects.

Estimates of the block or environmental effects are usually of less interest. It may be verified that the least squares estimate of β_j is

$$b_j = \frac{B_j}{12} - m$$

where B_j is the observed block total.

3.24 Tests of Significance and Confidence Limits. When the estimates of the treatment effects have been obtained, the details of the subsequent study of them vary with the type of experiment. Usually, however, some of the following procedures form the basis for the final conclusions: (i) a test of significance of the null hypothesis that the effects τ_i and τ_u of two treatments are identical, or, more generally, of the hypothesis that some linear combination of the τ_i is zero; (ii) the construction of confidence limits for the difference $(\tau_i - \tau_u)$ between the effects of two treatments, or for a linear function of the treatment effects; (iii) a test of significance of the null hypothesis that a group of treatments $\tau_1, \tau_2, \dots, \tau_p$ all have identical effects. In this section the theory governing these procedures is presented briefly, without proof. The arithmetical methods are described in the next section.

i. *Test of significance of the difference between two treatment effects.*
From equation (3.9) we have

$$t_i - t_u = (\tau_i - \tau_u) + (\bar{e}_i - \bar{e}_u) \quad (3.9)$$

Since the e 's are normally distributed with zero means, it follows by theory that $(t_i - t_u)$ is normally distributed about the true difference $(\tau_i - \tau_u)$. Further, the variance of $(t_i - t_u)$ may be shown to be

$$\sigma^2 \left(\frac{1}{r_i} + \frac{1}{r_u} \right)$$

where r_i and r_u are the numbers of replications of t_i and t_u , respectively. Since it is useful to be able to calculate the variance of any comparison in which we are interested, the rules for doing this are given in section 3.5.

In order to proceed we need an estimate of the experimental error variance σ^2 . From theory, the best estimate is known to be

$$s^2 = \sum \frac{(y_{ijk} - m - t_i - b_j)^2}{n_e}$$

where the residual sum of squares is taken over all observations. The divisor n_e , called the *number of degrees of freedom* (d.f.) in the estimated error, is given by the rule

$$n_e = (\text{Total number of observations}) - (\text{number of independent parameters that were estimated})$$

The word *independent* is introduced because we nearly always include in the original model more parameters than are strictly necessary. In the example we found that we could assume one linear relation (3.6) among the t 's and one (3.7) among the b 's. Consequently the number of independent parameters is

$$1 \text{ (for } m) + 8 \text{ (for the } t\text{'s)} + 3 \text{ (for the } b\text{'s)} = 12$$

Since there are 48 observations, n_e is 36.

Hence, if the null hypothesis is true, i.e., $\tau_i = \tau_u$, then $(t_i - t_u)$ is normally distributed with mean zero and estimated variance

$$s^2 \left(\frac{1}{r_i} + \frac{1}{r_u} \right)$$

The ratio

$$\frac{t_i - t_u}{s \sqrt{\frac{1}{r_i} + \frac{1}{r_u}}}$$

is known to follow Student's t -distribution with n_e , or 36 d.f. This is the quantity used for a test of significance of the null hypothesis.

With a more complicated function of the estimated treatment effects, say $L = (w_1 t_1 + w_2 t_2 + \dots + w_k t_k)$, where the w 's are any set of numbers, the procedure is essentially the same. The estimated variance of L is $s^2 f$, where f is a numerical factor found by the rules given in section 3.5. For a test of the null hypothesis that the true value of L is zero, we use the ratio $L/s\sqrt{f}$, which is distributed as Student's t with n_e degrees of freedom.

ii. *The construction of confidence limits.* The theory for a test of significance also leads to confidence limits for the unknown true difference $(\tau_i - \tau_u)$. For, when τ_i and τ_u are not necessarily equal, the quantity

$$\frac{(t_i - t_u) - (\tau_i - \tau_u)}{s \sqrt{\frac{1}{r_i} + \frac{1}{r_u}}}$$

may be shown to follow Student's t -distribution with n_e degrees of freedom. Hence the confidence limits are given by

$$(\tau_i - \tau_u) = (t_i - t_u) \pm sd(n_e, \alpha) \sqrt{\frac{1}{r_i} + \frac{1}{r_u}}$$

where $d(n_e, \alpha)$ is the value of Student's t corresponding to n_e degrees of freedom and the chosen confidence probability α . A similar method is used for the more general function L .

iii. *A test of the identity of a group of treatment effects.* Quite frequently the first test to be made is that of the hypothesis that all k treatments have produced identical effects, i.e., that all τ_i are equal. In other cases we are more interested in the hypothesis that some subgroup of the treatments, say treatments 1 to p , has produced the same effects. We consider the latter test, since it reduces to the first test if p is put equal to k .

This test is provided by a general theorem that has many applications. First, we find the least squares estimates of all treatment and environmental effects in the usual way, and compute the residual sum of squares

$$S_1^2 = \sum (y_{ijk} - m - t_i - b_j)^2$$

Next we start again and rewrite the mathematical model, inserting the restriction that $\tau_1 = \tau_2 = \dots = \tau_p$. That is, wherever any of these τ_i appears in the equation for an observation, we replace the τ_i by a common symbol, say τ' . The least squares estimates are computed for

this new set of equations, using the same observations. Again we compute the residual sum of squares which may be denoted by

$$S_2^2 = \sum (y_{ijk} - m' - t' - b_j')^2$$

This will always be found to be at least as large as S_1^2 . Finally, the theorem states that, if the null hypothesis is true, the quantity

$$\frac{S_2^2 - S_1^2}{p - 1} \div \frac{S_1^2}{n_e}$$

follows Snedecor's F -distribution with $(p - 1)$ and n_e degrees of freedom.

Although the procedure may appear rather complex, the test criterion used is a reasonable one. S_1^2 is the sum of squares of deviations of the observations from the values predicted for them by the original model, so that it measures how closely the original model agrees with the data. S_2^2 plays the same role for the restricted model. Consequently, if S_2^2 is much larger than S_1^2 , we are inclined to think that the restricted model does not fit the data nearly so well as the original model, and therefore to reject the null hypothesis that the restricted model is the correct one. However, the value of $(S_2^2 - S_1^2)$ alone does not provide a measure of the improvement in the fit with the original model. If, for instance, this difference is 10, then, other things being equal, we should regard the improvement in fit as greater when S_1^2 is 1 than when S_1^2 is 1000. It is this type of consideration that leads to the use of the ratio $(S_2^2 - S_1^2)/S_1^2$ as the essential part of the test criterion.

Although the theorem postulates two separate sets of least squares solutions, nearly all designs are constructed so that only one set of solutions must be found in practice. A number of designs which could have been included in this book were omitted because the least squares analysis seems too cumbersome for frequent use.

3.25 The Analysis of Variance. All the procedures (t -tests, F -tests, and construction of confidence limits) use the residual sum of squares, which will often be called the *error* sum of squares. This quantity could be found by calculating for each observation y_{ijk} the value $(m + t_i + b_j)$ predicted by the least squares solution. The sum of the squares of the differences between observed and predicted values could then be obtained. This method is slow, and the error sum of squares is much more quickly computed by a technique known as the *analysis of variance*.

In the original model, each observation is represented as the sum of four components due respectively to the general mean, the effect of the treatment, the environmental effect, and the residual effect. In the same

way the analysis of variance partitions the sum of squares of the observations into four sums of squares, one attributable to the general mean, one to differences between the estimated effects of the treatments, one to the environmental effects which the experiment is capable of measuring, and lastly one which is the residual or error sum of squares. In most cases we compute the original sum of squares and the first three components, obtaining the error sum of squares by subtraction.

The analysis of variance provides much more than a short-cut method of securing the error sum of squares. The sum of squares due to treatments is the quantity $(S_2^2 - S_1^2)$ needed for the F -test of the hypothesis that no differences exist between the effects of the treatments. By a slight extension the analysis also supplies the sum of squares required for testing the equality of the effects of a subgroup of the treatments. The component due to environmental effects enables us to estimate by how much the accuracy of the experiment has been increased by eliminating these effects from the estimates of the treatment means.

The analysis of variance depends on a number of algebraic relations which will be illustrated for the eelworm experiment. From section 3.23 it will be recalled that the least squares normal equation for any unknown, say t_i , was

$$\sum (y_{ijk} - m - t_i - b_j) = 0$$

summed over all observations which received the i th treatment. If we multiply this equation by t_i and add the equations for different treatments together, we obtain the equation

$$\sum t_i(y_{ijk} - m - t_i - b_j) = 0 \quad (3.10)$$

where the sum is now over *all* observations, since every observation is associated with one and only one t_i . Similarly we may establish the relations

$$\sum m(y_{ijk} - m - t_i - b_j) = 0 \quad (3.11)$$

$$\sum b_j(y_{ijk} - m - t_i - b_j) = 0 \quad (3.12)$$

where both sums extend over all observations.

A few more relations of this type are required. If we add the estimates t_i over all plots in the j th block, the sum is

$$(t_1 + t_2 + t_3 + \cdots + t_8) + 4t_9$$

But this is zero because of equation (3.6) in which a linear relation among the t 's was introduced. Hence, if we multiply by b_j and add over all blocks, we establish the relation

$$\sum b_j t_i = 0 \quad (3.13)$$

where the sum is again over all observations. By similar arguments we prove the additional relations

$$\sum mb_j = 0; \quad \sum mt_i = 0 \quad (3.14)$$

These relations lead to the partition of the sum of squares of the observations. Write

$$y_{ijk} = m + t_i + b_j + (y_{ijk} - m - t_i - b_j)$$

Square both sides, and add over all observations. The six relations (3.10) to (3.14) show that all six sums arising from cross-product terms on the right-hand side add to zero. Consequently, we have the following analysis of sums of squares.

$$\sum y_{ijk}^2 = \sum m^2 + \sum t_i^2 + \sum b_j^2 + \sum (y_{ijk} - m - t_i - b_j)^2 \quad (3.15)$$

This equation is the basis of the analysis of variance of the results.

3.26 Application to the Example. To obtain the left-hand side of (3.15) in practice, we simply compute the sum of the squares of all observations. For the eelworm data this will be found to be 5,481,198. The components due to the mean, to treatments, and to blocks are not usually calculated from (3.15) as it stands, because it is quicker and more accurate to obtain them from *totals* rather than from the estimated effects.

Thus, for the mean, the contribution is $48m^2$, since there are 48 observations. If G is the grand total of all observations, 14,680, this contribution may be written $G^2/48$, or $(14,680)^2/48$, which amounts to 4,489,633. This term is sometimes called the correction for the mean.

Written in full, the sum of squares for treatments is

$$4(t_1^2 + t_2^2 + \cdots + t_8^2) + 16t_9^2 \quad (3.16)$$

But from (3.8),

$$t_i = \frac{T_i}{4} - m \quad (i = 1, 2, \dots, 8); \quad t_9 = \frac{T_9}{16} - m$$

where T_i is the observed treatment total. By substitution, (3.16) becomes

$$\frac{1}{4}[(T_1 - 4m)^2 + (T_2 - 4m)^2 + \cdots + (T_8 - 4m)^2] + \frac{1}{16}(T_9 - 16m)^2$$

When each parenthesis is expanded, we have

$$\frac{T_1^2 + T_2^2 + \cdots + T_8^2}{4} + \frac{T_9^2}{16} - 2m(T_1 + T_2 + \cdots + T_8 + T_9) + 48m^2$$

Now the sum of the observed treatment totals T_i is the grand total G , or $48m$. Consequently the last two terms above may be amalgamated to give

$$\frac{T_1^2 + T_2^2 + \dots + T_8^2}{4} + \frac{T_9^2}{16} - 48m^2 \quad (3.17)$$

Note that the last term is the correction for the mean, already found to be 4,489,633. Expression (3.17) is an example of the more general expression

$$\sum_{i=1}^k \frac{T_i^2}{r_i} - C$$

which gives the treatments sum of squares when the i th treatment is replicated r_i times, C being the correction for the mean.

To obtain the numerical value we first calculate the treatment totals as shown in table 3.2.

TABLE 3.2 TREATMENT TOTALS FROM TABLE 3.1

Level of application	CN	CS	CM	CK	Totals
0			5858		5858
1	1066	928	1431	892	4317
2	1265	877	1241	1122	4505

The treatments sum of squares is given by

$$\frac{(1066)^2 + (1265)^2 + \dots + (1122)^2}{4} + \frac{(5858)^2}{16} - 4,489,633 = 157,448$$

The blocks sum of squares, $\sum b_j^2$, can likewise be expressed in terms of block totals as follows.

$$\begin{aligned} & \frac{B_1^2 + B_2^2 + B_3^2 + B_4^2}{12} - C \\ &= \frac{(4383)^2 + (3075)^2 + (4752)^2 + (2470)^2}{12} - 4,489,633 = 289,427 \end{aligned}$$

where the observed block totals have been inserted from table 3.1. The divisor 12 is the number of observations per block.

The complete analysis of variance is shown in table 3.3. It is not customary to display the correction term for the mean, since this is of no particular interest. The item labelled "total" at the foot of the table is the original sum of squares, *minus* the correction for the mean (5,481,198

$- 4,489,633 = 991,565$). This may be shown to be equal to the sum of squares of deviations of the observations from their mean; i.e.,

$$\sum y_{ijk}^2 - 48m^2 = \sum (y_{ijk} - m)^2$$

TABLE 3.3 ANALYSIS OF VARIANCE OF THE DATA IN TABLE 3.1

Source of variation	Degrees of freedom (d.f.)	Sum of squares (s.s.)	Mean square (m.s.)
Treatments	8	157,448	19,681
Blocks	3	289,427	96,476
Error (by subtraction)	36	544,690	15,130
Total	47	991,565	

The column which has been computed above is, of course, the column labelled "sum of squares." The other two columns remain to be explained. The "degrees of freedom" associated with any component are the number of independent parameters required to describe that component in the model. In the case of treatments, this always equals one less than the number of treatments, and similarly for blocks. For the total s.s., the number is the number of observations (48) less one representing the contribution of the mean.

The degrees of freedom have two principal uses. First, by subtraction, they give the degrees of freedom for error (36). As mentioned in section 3.24, this is the divisor needed for the error s.s. in order to estimate the error variance σ^2 . The estimate is $544,690/36$, or $15,130$, which is called the *error mean square* and is shown in the right-hand column of table 3.3. Second, the degrees of freedom for treatments are used in an F -test of the hypothesis that all k treatments produced the same effects. In section 3.24 the value of F for testing this hypothesis was given as

$$F = \frac{S_2^2 - S_1^2}{k - 1} \div \frac{S_1^2}{n_e}$$

The denominator S_1^2/n_e is the error m.s. In the numerator the quantity $(S_2^2 - S_1^2)$ may be shown to be the treatments s.s., while $(k - 1)$ is the number of degrees of freedom associated with treatments. Thus the numerator is $157,448/8$, or $19,681$, called the *treatments mean square*. The F -ratio for this experiment is $19,681/15,130$, or 1.30 , with 8 and 36 d.f.

In the same way the mean square for blocks forms the numerator for the F -test of the hypothesis that all block effects are identical. Although this test is rather seldom of interest, the mean square for blocks has other

uses. By formulae which will be given with the individual designs, this mean square leads to an estimate of the error variance that would have been obtained if the experiment had not been grouped into blocks.

With some designs the structure of the analysis of variance is more complicated than in the example above. What happens is that the treatment and environmental contributions to the sum of squares are entangled. If we calculate the treatments s.s., a different result is obtained according to whether we assume that block effects are present or absent, and vice versa for blocks. In such cases the usual procedure is to present the blocks s.s. as calculated when treatment effects are ignored, and the treatments s.s. as calculated when block effects are taken into account. These items are sometimes called "blocks, ignoring treatments" and "treatments, eliminating blocks," respectively. If calculations are made in this way, the two terms may still be added so as to obtain the error s.s. by subtraction; further, the treatments s.s. is still the appropriate one for an F -test of the treatment effects.

This presentation of the analysis of variance is necessarily very inadequate. For further reading, reference should be made to Fisher (3.2, 3.3) and Snedecor (3.4), whose discussions involve little mathematics. For presentation of points of theory, see Kendall's book (3.5) and papers by Fisher (3.6, 3.7) and Yates (3.8). Unfortunately, no single reference contains a really comprehensive account of the subject.

3.3 Accuracy in Computations

3.31 Original Records. If there exists a series of rules which will guarantee that all computations are accurately done, we do not know them. The following notes may help in maintaining a high standard of accuracy.

The first place where errors may occur is in the original records. These should be made in a clearly legible and permanent form. Where scale readings are being taken, the habit of checking each reading immediately after it has been written down is a useful one. As the observations are being recorded, any that appear anomalous should be examined and the reason sought. Gross errors are often eliminated in this way. If on checking such an observation no error or explanation is found, a note to this effect should be made. Such notes are helpful in situations where the later analysis is done by a different person. Most statisticians who analyze other people's experiments have encountered the perplexing problem where an observation is so completely out of line that a gross error is suspected, yet the original records contain no comment about

the observation, and the recorder's memory of the circumstances in which it was taken has faded.

Copying from the original records to a form suitable for analysis may also introduce errors. Since cross-checking is rather a soporific task, mistakes tend to be overlooked even when two persons are used for the check. Sometimes foresight can eliminate the need for copying.

3.32 Checks in the Analysis of Variance. The analysis of variance itself consists of two sorts of operations. First, a number of totals are formed. These are nearly all self-checking, for the fact that the treatment totals and the block totals both add to the grand total may be accepted as proof of their correctness. This is not a watertight check, since compensating errors in two treatment totals pass undetected. Also it does not ensure that treatments and blocks are correctly identified, and cases have occurred where treatment totals were mistakenly labelled as block totals, and vice versa. But apart from such rare events the sum check is as satisfactory as any.

The second stage, the computation of the various sums of squares, is not self-checking and must be checked by recomputation. The most accurate method is to have two computers carry out the analysis independently, comparing results only when both have finished the task. The practice of having one computer "check" the results of the other, though more convenient, always seems to produce a certain number of errors. This is to be expected, because, when the first calculations are nearly always correct, as they should be, it is difficult for the checker to avoid the presupposition that they are correct, with resulting mental laziness. Recomputation by the same person, though often unavoidable, is probably less accurate than the two preceding methods.

An expert who wishes to do so can make the analysis self-checking. For instance, the error s.s. can be computed independently of the other sums of squares, sometimes quite easily and sometimes with more difficulty. If this sum of squares agrees with that found by subtraction, all the items in the "sums of squares" column may be regarded as correct, apart possibly from errors in labelling. However, with the computational methods that we present, checking is required except in a few cases specifically noted.

The steps in the preparation of summary tables (e.g., calculation of treatment means from totals) are usually partly but not entirely self-checking. For example, the fact that treatment means average to the general mean forms a check. Care should be taken to recompute those steps that are not self-checked.

With the more complex designs, where treatment means have to be adjusted in order to eliminate the environmental effects, errors have arisen because the computing instructions were misunderstood. For instance, the adjustments made were ten times as large as they should have been. A person thoroughly familiar with the design would have known that the adjustments were ridiculously large, but someone new to the design may have no idea what size of adjustment it is reasonable to expect. This danger with complex designs should not be ignored. It can be minimized by careful study of the method of analysis for any design that it is proposed to use.

3.33 Number of Figures to be Retained. Since the primary purpose of the analysis of variance is to obtain an accurate value for the error m.s., the number of figures which it is worth retaining during the calculations depends on the size of this figure. This is not known in advance, but often a rough idea of the coefficient of variation (ratio of the standard error to the mean of the experiment) is available from previous experience. For the *original data* from which the analysis of variance is computed, a crude rule which errs on the safe side is as follows. Record the original data to 4 significant figures if the coefficient of variation is between 0.4 and 4%, to 3 if it is between 4 and 40%, and to 2 if it exceeds 40%. Let us apply this rule to the eelworm data. Since experiments on eelworms are not very common, there is not much basis on which to predict the coefficient of variation. Suppose that we guess that it is unlikely to be below 20% (it actually turned out to be 40%). Then, for safety, data should be recorded to 3 significant figures. Since the average eelworm number per plot is around 300, this means that data are recorded to the nearest eelworm, which is as accurately as they could be recorded.

A more precise rule, requiring a little calculation, is that the rounding interval should not exceed one-quarter of the standard error per observation. To apply this rule to the eelworm data, starting with the same premise that the coefficient of variation is 20%, we note that the mean per plot in the experiment is about 300. Thus the predicted value of the standard error per observation is about 60, so that the rounding interval should not exceed 15. It will be quite satisfactory to round each observation to the nearest 10 eelworms. The discrepancy from the result given by the previous rule illustrates the fact that the first rule tends to be too conservative.

The advisability of deliberately rounding records already taken should be left to the judgment of the experimenter. With modern calculating machines it may be more expeditious to carry out the computations with

a few unnecessary figures than to take time to perform and check the rounding. The rules are most likely to be useful where computing devices are poor or unavailable.

In the analysis of variance itself, it is generally as well to carry the full number of figures obtained from the uncorrected sum of squares; e.g., if the original data contain one decimal place, the sum of squares and hence the analysis of variance will contain two decimal places. This number will usually be excessive, but no more time is likely to be wasted in writing down the unnecessary digits than in estimating how many digits should be retained.

In the final presentation of results, on the other hand, superfluous digits should be strictly avoided. They make the conclusions more difficult to grasp, impede rapid mental comparisons, and give some readers an erroneous impression of the accuracy of the results. A treatment mean should be rounded to one-tenth of its estimated standard error; that is, if the estimated standard error of the treatment means is 2.56, the means could be rounded to 0.256 and should be rounded to 0.1, i.e., to one decimal place, since this is the nearest rounding interval that is convenient. In some experiments (e.g., those of the split-plot type) different treatment comparisons have different standard errors. Here the rounding interval should be decided from those comparisons that have the smallest error.

3.34 Identification of Data. The practice should be followed of labeling all data, from original records to summary tables, so that another person at a later time can tell what the data mean. Much data, expensive to collect and potentially valuable for some research, have had to be discarded or destroyed because the original collector cannot be reached, and new investigators are unable to discover what measurements were taken and under what circumstances. Sometimes the units in which the observations or summary tables are presented cannot be found, or the crop on which a field experiment was carried out, or the nature of the experimental treatments. Frequently, data serve their most fruitful purpose when some later investigator gathers from many places all the material bearing on some question, and it should be a habit to facilitate rather than hinder such research.

3.4 Subdivision of the Sum of Squares for Treatments

3.41 Reasons for Subdivision. Often an experiment is planned to provide the answers to a number of different questions, not necessarily connected with one another. The eelworm experiment is a partial

though not an ideal example. One way in which the questions asked might be framed is as follows. (i) Is soil fumigation effective? Here we are interested in the average reduction in eelworm numbers taken over all fumigants. (ii) Are there differences in the effectiveness of different fumigants? For this question, comparisons of the reductions due to different fumigants are relevant. (iii) What is the relative effectiveness of single and double doses? Since the simplest reasonable hypothesis is that the reduction in eelworm numbers to the double dose is twice that to the single dose, we might narrow this question by asking: Is the average response to fumigation proportional to the amount of dressing, or in other words is the average response curve linear? (iv) Finally, if there is some indication of curvature in the responses, we might ask whether the amount of curvature is the same for all fumigants.

With experiments of this type the F -test of the complete treatments m.s. is not particularly helpful because it is directed, as it were, at a mixture of several diverse questions. By an extension of the analysis of variance, we can subdivide the treatments s.s. into a number of components that are more relevant to the individual questions. Moreover, an F -test can be made on the mean square for each component. The rules for subdivision are given in succeeding sections. Effective use of the device requires long practice applied to a considerable number of experiments.

3.42 Subdivision into Single Components. The simplest case, where all treatments have the same number r of replications, is considered first. The calculations are best made from the treatment totals T_i . Examples of the sort of quantity that we are interested in studying are

$$T_1 - T_2; \quad \frac{T_1 + T_2}{2} - T_3; \quad T_1 + T_2 - (T_3 + T_4)$$

Quantities of this type are called *linear functions* of the T 's. Note that the sum of the coefficients of the T 's is always zero, as it must be if the quantity is to represent a comparison among the T 's. These ideas may be formalized as follows.

Definition. Any linear function

$$z_w = l_{w1}T_1 + l_{w2}T_2 + \cdots + l_{wk}T_k$$

is called a *comparison* among the T 's if

$$l_{w1} + l_{w2} + \cdots + l_{wk} = 0$$

Rule 1. If z_w is any comparison among the T 's, the quantity

$$\frac{z_w^2}{D_w}, \text{ where } D_w = r(l_{w1}^2 + l_{w2}^2 + \cdots + l_{wk}^2)$$

is a component of the sum of squares for treatments and represents 1 d.f.

This rule gives the divisor D_w to be used so that the comparison z_w may be included as a part of the treatments s.s. in the analysis of variance. Incidentally, the rule implies that z_w^2/D_w cannot exceed the treatments s.s. If it is found to do so, this is a sure sign of a mistake in computation.

Definition. Two comparisons z_1 and z_2 are said to be *orthogonal* if

$$l_{11}l_{21} + l_{12}l_{22} + \cdots + l_{1k}l_{2k} = 0$$

This "sum of products" relation is very important. By theory, it ensures that z_1 and z_2 are distributed independently of each other. Its relevance to the partition of the treatments s.s. appears in rule 2.

Rule 2. If z_1 and z_2 are orthogonal, then z_2^2/D_2 is a component of

$$\text{Treatments s.s.} - \frac{z_1^2}{D_1}$$

This means that, if we divide the treatments s.s. into the contribution from a comparison z_1 and the remainder, and now wish to subdivide the remainder, we must choose comparisons that are orthogonal to z_1 . Similarly, after removing the contribution of z_2 , the next comparison z_3 must be orthogonal to both z_1 and z_2 , and so on. This leads to the next rule.

Rule 3. If the comparisons $z_1, z_2, \cdots, z_{k-1}$ are mutually orthogonal, i.e., every pair is orthogonal, then

$$\text{Treatments s.s.} = \frac{z_1^2}{D_1} + \frac{z_2^2}{D_2} + \cdots + \frac{z_{k-1}^2}{D_{k-1}}$$

This algebraic identity partitions the treatments s.s. [which has $(k - 1)$ degrees of freedom] into $(k - 1)$ components, each representing a single degree of freedom.

Starting with a specified z_1 , we can always find $z_2, z_3, \cdots, z_{k-1}$ so as to construct a complete orthogonal set. In fact, it may be proved that there is considerable freedom of choice in this process. When all but the final z_{k-1} have been selected, there is only one possible choice for z_{k-1} , but at any previous stage many z 's can be found that are orthogonal to all preceding z 's. The experimenter must choose the z 's that are most relevant for purposes of interpretation.

Table 3.4 shows the single components most frequently used for interpretative purposes when there are three or four treatments. It is as-

TABLE 3.4 SAMPLE SETS OF SINGLE COMPONENTS

Three treatments

i. Equally spaced increments of one ingredient				ii. Two qualities of an ingredient and a control			
	z_1	z_2			z_1	z_2	
T_1	-1	1		T_1 (0)	-2	0	
T_2	0	-2		T_2 (a_1)	1	-1	
T_3	1	1		T_3 (a_2)	1	1	
Component	Lin.	Quad.			Effect	Quality	
divisor D	$2r^*$	$6r$		Component	of a	diff.	
				divisor D	$6r$	$2r$	

Four treatments

i. Equally spaced increments of one ingredient				ii. Two comparable types of ingredient, a and b , and two qualities of each type			
	z_1	z_2	z_3		z_1	z_2	z_3
T_1	-3	+1	-1	T_1 (a_1)	+1	+1	0
T_2	-1	-1	+3	T_2 (a_2)	+1	-1	0
T_3	+1	-1	-3	T_3 (b_1)	-1	0	+1
T_4	+3	+1	+1	T_4 (b_2)	-1	0	-1
Component	Lin.	Quad.	Cubic			Quality	Quality
D	$20r$	$4r$	$20r$	Component a vs. b		diff. within a	diff. within b
				D	$4r$	$2r$	$2r$

iii. Different levels of each of two different ingredients a and b

		z_1	z_2	z_3	
T_1	(a_1b_1)	-1	-1	+1	
T_2	(a_2b_1)	+1	-1	-1	
T_3	(a_1b_2)	-1	+1	-1	
T_4	(a_2b_2)	+1	+1	+1	
Component		Average response to a	Average response to b	Interaction	
D		$4r$	$4r$	$4r$	

r^* = number of replicates, assumed the same for all treatments. The coefficients given above are not valid if r varies from treatment to treatment.

sumed that all treatments have the same number of replicates. The divisors D are those required for inserting the square of z in the analysis

of variance. The reader should verify that these divisors satisfy rule 1, and that each set of z 's forms a complete orthogonal set.

With three treatments the two sets presented are essentially the same, though the meaning attached to each z is different. Case i applies in an experiment where we have, say, zero, single, and double dressings of a fumigant, and wish to isolate the linear and quadratic components of the response curve. The response found by fitting a straight line to the three levels is represented by z_1 , while z_2 measures the deviation from a linear response. Case ii arises when we have comparable dressings of two different fumigants, and wish to examine the average effect of fumigation (z_1) and the difference between the fumigants (z_2).

With four treatments, case i shows zero, single, double, and triple applications of the same ingredient; here the response curve can be divided into its linear, quadratic, and cubic components. Alternatively, in case ii, we might be testing the wearing qualities of two types of "100% wool" suits and two types of suits made of wool-rayon mixtures. Case iii is used, for instance, when we examine the effects of two different amounts of sugar and two different amounts of vanilla in the preparation of a cake. Experiments of this type are discussed more fully in chapter 5.

When different treatments have differing numbers of replicates, the rules are changed slightly. If the i th treatment has r_i replicates, we have shown that the treatments s.s. is

$$\frac{T_1^2}{r_1} + \frac{T_2^2}{r_2} + \cdots + \frac{T_k^2}{r_k} - \frac{(T_1 + T_2 + \cdots + T_k)^2}{(\sum r_i)}$$

The changes to be noted are:

i. A linear function

$$z_w = l_{w1}T_1 + l_{w2}T_2 + \cdots + l_{wk}T_k$$

is a comparison among the treatment totals T_i if

$$r_1l_{w1} + r_2l_{w2} + \cdots + r_kl_{wk} = 0$$

ii. The divisor required for z_w^2 is

$$D_w = r_1l_{w1}^2 + r_2l_{w2}^2 + \cdots + r_kl_{wk}^2$$

iii. Two comparisons z_1 and z_2 are orthogonal if

$$r_1l_{11}l_{21} + r_2l_{12}l_{22} + \cdots + r_kl_{1k}l_{2k} = 0$$

3.43 Incomplete Subdivisions. The subdivision need not be complete in the sense that it is composed entirely of single components. In the eelworm experiment we suggested that the treatments s.s., with 8

components, might be divided into the following parts: (i) a single component representing the average response to fumigation; (ii) a single component representing the deviation of the average response from linearity; (iii) a part representing differences in the responses to the different fumigants. Since there are 4 fumigants, this part comprises 3 components; (iv) a part representing differences in the curvature of the response curves for different fumigants: this also has 3 components. Some rules for partial subdivision will be given and applied to this example. The most general rule is as follows.

Rule 4. A set of g quantities Q_i are independent components of the treatments s.s., with n_i degrees of freedom respectively, if

$$Q_i = \frac{z_{i1}^2}{D_{i1}} + \frac{z_{i2}^2}{D_{i2}} + \cdots + \frac{z_{in_i}^2}{D_{in_i}} \quad (i = 1, 2, \dots, g)$$

where all z_{ij} are mutually orthogonal comparisons, and the D_{ij} are the appropriate divisors. Further, the remainder

$$\text{Treatments s.s.} - \sum Q_i$$

is an independent component of the treatments s.s. with

$$[(k-1) - \sum n_i] \text{ degrees of freedom}$$

This rule, which is a deduction from previous rules, is not very helpful to the beginner. With experience it often becomes easy, without carrying out the details, to see whether a set of Q_i can be expressed in this way. Some more specialized rules follow.

Rule 4a. If T_1, T_2, \dots, T_p are the totals for any set of treatments, then

$$Q = \frac{T_1^2}{r_1} + \frac{T_2^2}{r_2} + \cdots + \frac{T_p^2}{r_p} - \frac{(T_1 + T_2 + \cdots + T_p)^2}{\sum r_i}$$

is a component of the treatments s.s., with $(p-1)$ degrees of freedom.

This rule states that the sum of squares of deviations among any subgroup of the treatments, with proper divisors, is part of the treatments s.s.

Rule 4b. Sometimes the treatments (or part of them) can be divided into a number of groups. The number of treatments in a group need not be constant, and different treatments may have different amounts of replication. Let T_{ij} be the total for the j th treatment in the i th group, and let

$$S_i = T_{i1} + T_{i2} + \cdots + T_{ip_i};$$

$$R_i = r_{i1} + r_{i2} + \cdots + r_{ip_i} \quad (i = 1, 2, \dots, g)$$

Then let

$$Q_i = \frac{T_{i1}^2}{r_{i1}} + \frac{T_{i2}^2}{r_{i2}} + \cdots + \frac{T_{ip_i}^2}{r_{ip_i}} - \frac{S_i^2}{R_i} \quad (i = 1, 2, \dots, g)$$

$$Q_{g+1} = \frac{S_1^2}{R_1} + \cdots + \frac{S_g^2}{R_g} - \frac{(S_1 + \cdots + S_g)^2}{R_1 + \cdots + R_g}$$

In this case Q_i represents the sum of squares of deviations among the treatments in the i th group, with $(p_i - 1)$ degrees of freedom, while Q_{g+1} is the sum of squares of deviations among the group totals with $(g - 1)$ degrees of freedom. These Q 's are independent components of the treatments s.s.

Rule 4c. This applies when all treatments have the same number r of replicates, and every group contains p treatments. Let

$$z_i = l_1 T_{i1} + l_2 T_{i2} + \cdots + l_p T_{ip}$$

where the set of l 's remains the same in all groups. Then

$$Q = \sum \frac{(z_i - \bar{z})^2}{r \sum l_i^2}$$

is a component of the treatments s.s., with $(g - 1)$ degrees of freedom. This rule is useful when we wish to compare linear functions of the treatments within each group. Note that the divisor in Q is the same as the divisor that would be used for a single z_i .

Applications. In the celworm experiment, the *average* effects of fumigation are obtained from a comparison of the total for the control with the totals for the single and double levels of fumigation. Consequently, we might regard the 9 treatments as divided into 3 groups: the control, the single levels, and the double levels. By rule 4b, this gives the following partition of the treatments s.s. (the treatment totals are found in table 3.2, p. 50).

Between fumigants (single level):

(3 d.f.)

$$\frac{(1066)^2 + (928)^2 + (1431)^2 + (892)^2}{4} - \frac{(4317)^2}{16} = 45,461$$

Between fumigants (double level):

(3 d.f.)

$$\frac{(1265)^2 + (877)^2 + (1241)^2 + (1122)^2}{4} - \frac{(4505)^2}{16} = 23,641$$

Between levels:

(2 d.f.)

$$\frac{(5858)^2 + (4317)^2 + (4505)^2}{16} - \frac{(14,680)^2}{48} = 88,347$$

The three sums of squares add to 157,449, in agreement with the treatments s.s., 157,448, as given in table 3.3, p. 51.

The next step is to divide the sum of squares between levels into a component representing the average response to fumigation and one representing the deviation from a linear response. By least squares theory, the former is given by a comparison of the double level with the zero level. That is, we may take

$$z_1 = S_2 - S_0 = 4505 - 5858 = -1353$$

where the subscript refers to the level. The deviation from linearity is measured by the orthogonal comparison

$$\begin{aligned} z_2 &= (S_2 - S_1) - (S_1 - S_0) = S_2 + S_0 - 2S_1 \\ &= 4505 + 5858 - 2(4317) = 1729 \end{aligned}$$

Since each S is a total over 16 plots, the divisors are 32 and 96 respectively. Hence the contributions to the sum of squares are:

$$\text{Average linear response: } \frac{(1353)^2}{32} = 57,207$$

$$\text{Average curvature: } \frac{(1729)^2}{96} = 31,140$$

The subdivision found thus far might be presented as shown below.

	d.f.	s.s.	m.s.
Between treatments	8	157,449	19,681
Average linear response	1	57,207	57,207
Average curvature	1	31,140	31,140
Between single levels of fumigants	3	45,461	15,154
Between double levels of fumigants	3	23,641	7,880

This is not the subdivision envisaged at the beginning of this section. Instead, it was proposed to divide the final 6 d.f. into 3 representing differences in the responses to individual fumigants and 3 representing differences in the curvatures of the individual response curves. This separation is more difficult.

Consider a comparison of (CN) and (CS). Estimates of the difference between these fumigants are available both at the single (1CN) and the double (2CN) level. If the effects of both fumigants are proportional to the amounts of dressing, the true difference at the double level will be twice that at the single level. On this assumption, two independent estimates of the difference between (CN) and (CS), for a single dressing, are

$$[(1CN) - (1CS)] \quad \text{and} \quad \frac{[(2CN) - (2CS)]}{2}$$

The second estimate is considerably more accurate than the first, since its variance is only $\frac{1}{4}$ as large. Statistical theory shows that such estimates are combined by weighting each inversely as its variance.

Hence, on the assumption of linearity, the most accurate estimate of the difference per unit dressing is

$$\frac{[(1\text{CN}) - (1\text{CS})] + (4)(\frac{1}{2})[(2\text{CN}) - (2\text{CS})]}{1 + 4} = \frac{[(1\text{CN}) - (1\text{CS})] + 2[(2\text{CN}) - (2\text{CS})]}{5} \quad (3.18)$$

This shows that differences in the linear responses to the 4 fumigants are measured by comparisons of the quantities $(1\text{CN}) + 2(2\text{CN})$, etc. These quantities are given below.

(Single level) + 2(double level)				
(CN)	(CS)	(CM)	(CK)	Total
3596	2682	3913	3136	13,327

It is for quantities of this kind that rule 4c is useful. This rule shows that the sum of squares of deviations of these quantities, when divided by $r \sum l_i^2$, or $(4) [(1)^2 + (2)^2]$, i.e., 20, is a component of the treatments s.s. This gives 43,408, representing differences in linear responses to the fumigants.

We now consider the final three components. The curvature of an individual response curve is measured by a comparison of the type

$$[(2\text{CN}) - (1\text{CN})] - [(1\text{CN}) - (0)] = [(2\text{CN}) + (0) - 2(1\text{CN})]$$

where (0) represents the total number of eelworms over 4 plots for the control treatment. When the difference between two fumigants is taken, the (0) term disappears and we obtain the comparison

$$[(2\text{CN}) - 2(1\text{CN})] - [(2\text{CS}) - 2(1\text{CS})] \quad (3.19)$$

It is easy to verify that this is orthogonal with comparisons like (3.18). Consequently differences in curvature are compared by means of the quantities shown below.

2(Single level) - (double level)				
(CN)	(CS)	(CM)	(CK)	Total
867	979	1621	662	4129

Rule 4c again applies. The divisor is 20, and the sum of squares contributes 25,693. The final separation is given in table 3.5. It is evident

from F -tests that none of the mean squares is significant at the 5% level, and only that for the average linear response approaches near to the 5% level. These results must not be taken as final conclusions for this experiment. As will be seen in section 3.8, additional data were recorded in this experiment which permit a more accurate analysis.

TABLE 3.5 SUBDIVISION OF THE TREATMENTS S.S. FOR THE EELWORM DATA

Source of variation	d.f.	s.s.	m.s.
Average linear response	1	57,207	57,207
Average curvature	1	31,140	31,140
Differences in linear response	3	43,408	14,469
Differences in curvature	3	25,693	8,564
Error	36	544,690	15,130

A more extensive discussion of this general topic, with a number of examples, is given in Snedecor's book (3.4), chapter 15. It is worth repeating that the amount and type of subdivision that should be done depend on the experiment, and to some extent on individual taste. As will be seen in the next section, any single component can be tested by means of a t -test derived from the treatment means, and some workers prefer to make the test in this way.

3.5 Calculation of Standard Errors for Comparisons among Treatment Means

3.51 Rules. We first give a rule (rule 5) that is more general than we need, because it is sometimes useful for other purposes.

Rule 5. The standard error of any linear function

$$z = l_1 y_1 + l_2 y_2 + \cdots + l_N y_N$$

of the individual observations is

$$\sigma_z = \sigma \sqrt{l_1^2 + l_2^2 + \cdots + l_N^2}$$

The *estimated* standard error is obtained by substituting s for σ , where s is the square root of the error m.s. in the analysis of variance. This rule supplies the standard error of any kind of linear function, provided that it has been expressed in terms of the *individual observations*. For linear functions of the *treatment means*, one of two special cases of the rule is used.

Rule 5a. If all treatments have the same number r of replications, the standard error of a linear function

$$z = l_1 \bar{y}_1 + l_2 \bar{y}_2 + \cdots + l_k \bar{y}_k$$

of the treatment means \bar{y}_i is

$$\sigma_z = \frac{\sigma}{\sqrt{r}} \sqrt{l_1^2 + l_2^2 + \cdots + l_k^2}$$

Rule 5b. If the i th treatment has r_i replications, the standard error of z is

$$\sigma_z = \sigma \sqrt{\frac{l_1^2}{r_1} + \frac{l_2^2}{r_2} + \cdots + \frac{l_k^2}{r_k}}$$

As in the general case, we substitute s for σ in order to obtain the estimated standard error. Finally, a rule that may save labor in complex cases is as follows.

Rule 6. If the linear functions z_1, z_2, \dots, z_p are mutually orthogonal, the standard error of

$$z = l_1 z_1 + l_2 z_2 + \cdots + l_p z_p$$

is

$$\sigma_z = \sqrt{l_1^2 \sigma_1^2 + l_2^2 \sigma_2^2 + \cdots + l_p^2 \sigma_p^2}$$

where σ_i^2 is the variance of z_i .

This rule enables the work to be done in two stages. Sometimes, in a large experiment, it is relatively easy to express the function z in terms of a number of familiar orthogonal functions, but rather tedious to write z in terms of original observations. The variances of the z_i may be found by rule 5 and substituted in rule 6. Note that different z_i may involve the same set of observations, provided that the z_i are orthogonal.

3.52 Examples. These rules will now be illustrated by application to the eelworm experiment. The treatment means are shown in table 3.6, the figures in parentheses denoting the order in which the treatments are numbered.

TABLE 3.6 MEAN NUMBERS OF EELWORMS PER PLOT

Level	Fumigant				Means
	(CN)	(CS)	(CM)	(CK)	
0			366(9)		366
1	266(5)	232(6)	358(7)	223(8)	270
2	316(1)	219(2)	310(3)	280(4)	281

From table 3.5 the standard error per plot is $\sqrt{(15,130)}$, or $s = 123$, with 36 d.f.

Example 1. Standard error of the average linear response. This is measured by the average response to the double dressing, i.e.,

$$z = \frac{1}{4}(\bar{y}_1 + \bar{y}_2 + \bar{y}_3 + \bar{y}_4) - \bar{y}_9 = -85$$

Since the first 4 means are based on 4 replicates, while y_9 has 16 replicates, we apply rule 5b, which gives

$$\begin{aligned} s_z &= (123) \sqrt{\left(\frac{1}{16}\right)\left(\frac{1}{4}\right) + \left(\frac{1}{16}\right)\left(\frac{1}{4}\right) + \left(\frac{1}{16}\right)\left(\frac{1}{4}\right) + \left(\frac{1}{16}\right)\left(\frac{1}{4}\right) + (1)\left(\frac{1}{16}\right)} \\ &= \frac{123}{\sqrt{8}} = 43.5 \end{aligned}$$

The value of Student's t is $85/43.5$, or 1.95.

As mentioned previously, the t -test of any comparison, as made above, is identical with the F -test of the corresponding component in the analysis of variance. The basic relationship is $F = t^2$; that is, if the 5% values of t for n degrees of freedom are read from the table and squared, their squares are the 5% F values for 1 and n degrees of freedom, and similarly for any other significance level. This result may be verified in the present instance. The value of t^2 is 3.80, with 36 d.f. From the analysis of variance in table 3.5, the F value for the average linear response is $57,207/15,130$ or 3.78, with 1 and 36 d.f., the two values agreeing apart from rounding errors. The reader may check that the same agreement holds for the average curvature.

Example 2. Response to the single dressing of (CN). The response is measured by

$$z = \bar{y}_5 - \bar{y}_9$$

Once again rule 5b is appropriate.

$$s_z = 123 \sqrt{\frac{1}{4} + \frac{1}{16}} = \frac{123\sqrt{5}}{4} = 68.8$$

Example 3. Difference between the linear responses to (CN) and (CS). In section 3.43 it was shown that this difference is

$$z = \frac{[\bar{y}_5 - \bar{y}_6 + 2\bar{y}_1 - 2\bar{y}_2]}{5}$$

Since all means have 4 replicates, rule 5a may be used.

$$s_z = \frac{123}{(2)(5)} \sqrt{1 + 1 + 4 + 4} = (12.3)(\sqrt{10}) = 38.9$$

Example 4. Although the experiment is not complex enough to exhibit a profitable application of rule 6, this example shows that the rule agrees with the other rules. Suppose that at the double level we had compared (CN) with (CS), and also the two chemicals with the two proprietary mixtures. The comparisons are

$$z_1 = \bar{y}_1 - \bar{y}_2; \quad z_2 = \frac{\bar{y}_1 + \bar{y}_2}{2} - \frac{\bar{y}_3 + \bar{y}_4}{2}$$

By rule 5a their standard errors are

$$\sigma_{z_1} = \frac{\sigma}{2} \sqrt{2}; \quad \sigma_{z_2} = \frac{\sigma}{2}$$

and it will be noted that the two functions are orthogonal.

If now we wished to compare the double dressing of (CN) with the mean of the 3 other double dressings, the comparison would be

$$z = \bar{y}_1 - \frac{\bar{y}_2 + \bar{y}_3 + \bar{y}_4}{3}$$

By a direct use of rule 5a, we have

$$\sigma_z = \frac{\sigma}{2} \sqrt{1 + \frac{1}{9} + \frac{1}{9} + \frac{1}{9}} = \frac{\sigma}{\sqrt{3}}$$

But alternatively we may write

$$z = \frac{2z_1}{3} + \frac{2z_2}{3}$$

and apply rule 6 to give

$$\sigma_z = \sqrt{\frac{4}{9} \sigma_{z_1}^2 + \frac{4}{9} \sigma_{z_2}^2} = \sigma \sqrt{\left(\frac{4}{9}\right) \left(\frac{1}{2}\right) + \left(\frac{4}{9}\right) \left(\frac{1}{4}\right)} = \frac{\sigma}{\sqrt{3}}$$

in agreement with the more direct method.

3.53 Testing Effects Suggested by the Data. In order that *F*- and *t*-tests be valid, the tests to be made in an experiment should be chosen before the results have been inspected. The reason for this is not hard to see. If tests are selected *after* inspection of the data, there is a natural tendency to select comparisons that appear to give large differences. Now large apparent differences may arise either because there are large real effects, or because of a fortuitous combination of the experimental errors. Consequently, in so far as differences are selected just because they seem to be large, it is likely that an undue proportion of the cases

selected will be those where the errors have combined to make the differences large. The extreme case most commonly cited is that of the experimenter who always tests, by an ordinary t -test, the difference between the highest and the lowest treatment means. If the number of treatments is large, this difference will be substantial even when the treatments produce no real differences in effect. It may be shown that with 3 treatments the observed value of t will exceed the 5% level in the table about 13% of the time. With 6 treatments the figure is 40%, with 10 treatments 60%, and with 20 treatments 90%. When the experimenter thinks that he is making a t -test at the 5% level, he is actually testing at the 13% level, or the 40% level, and so on. To summarize, the selection of those differences that look large and therefore "interesting" invalidates the ordinary tests of significance. The effect is to obtain too many significant results, or to raise the significance level of the test from the presumed 5% to some higher level, usually unknown.

On the other hand, the rule that no tests must be constructed after seeing the data seems contrary to sound scientific practice. Often the initial experiments in a line of research are conducted in order to "see what happens"; the research worker does not know in advance what comparisons he may wish to test. And even where hypotheses about the nature of the results can be set up, experiments often indicate strongly, as is later confirmed, that these hypotheses are erroneous and that the method of analysis based on them is inappropriate. Consequently, although the effects of selection cannot be ignored, they should not deter the experimenter from the most careful examination of his data. Any difference that is of interest, whether anticipated or not, should be tested. If the ordinary tests show that the difference is *not* significant, then at least we know that the difference can be accounted for without supposing any real effect. If an unexpected difference is statistically significant, the possibility of a selection effect should be borne in mind. When the point at issue is important, the best procedure is to conduct a new experiment specifically designed to confirm or disprove the indications from the previous results.

The reporting of such effects presents difficulties. The following is a suggestion. "The experiment also indicated an unexpected effect of Although this was statistically significant at the 5% level, it is contrary to previous experience, and must at present be regarded with caution. Further work on this point will be undertaken." The difficulties with this kind of writing are that we do not wish to imply that the effect is definitely established, and yet the writing should not encourage the reader to ignore any result that is contrary to our preconceived notions.

3.6 Subdivision of the Sum of Squares for Error

3.61 Reasons for Subdivision. The sum of squares for error can be partitioned into components in the same way as that for treatments. Although such subdivisions are not so frequently required as with treatments, they have a number of uses. Sometimes there is reason to believe that the error s.s. is not homogeneous; that is, the residual errors e do not all have the same variance σ^2 as postulated in the mathematical model. In this case, as will be shown in section 3.63, subdivisions of the error may be necessary in order to obtain valid t -tests. Occasionally a gross error in some observation may be suspected, and it is helpful to calculate the contribution of this observation to the error s.s. In addition, a subdivision may help in understanding the nature of the error m.s.

3.62 Rules for Subdivision. These follow the same general pattern as with treatments, though they are a trifle more complicated. It is necessary to go back to the normal equations (3.4) from which the treatment and block effects were estimated. These equations took the form

$$\sum (y_{ijk} - m - t_i - b_j) = 0 \quad (3.4)$$

In other words, the residuals that provide the estimates of error must add to zero over any treatment or over any block. Consequently, if the linear function

$$z = \sum l_{ijk} y_{ijk}$$

is to be a component of the error, the coefficients l_{ijk} must sum to zero over all observations that receive any specified treatment and also over all that are in any specified block. These are the tests by which we tell whether any proposed z is part of the error.

For any z which satisfies these tests, the contribution to the error s.s. is z^2/D , where by rule 5

$$D = \sum l_{ijk}^2$$

We now examine some components of the error s.s. in the eelworm experiment. Obviously, the difference z between the eelworm numbers on any two control plots in the same block satisfies the conditions. Similarly *any* comparison among the 4 control observations in a block is a component of the error. It follows that the sum of squares of deviations of these 4 observations from their mean contributes 3 d.f. to the error s.s. The 4 blocks together contribute 12 d.f. of this type of error. The sum of squares will be found to be 307,312. This component is a measure of the amount of variation among observations that receive the same

treatment and lie in the same block, and is the type of component that we should expect to constitute the error. However, this type of component is encountered only when some treatments have been replicated *within* the block.

Consider now the difference between two other treatments, say (1CN) and (1CS). Let

$$z_1 = (1CN)_1 - (1CS)_1$$

where the subscript 1 denotes that the two observations come from block 1. This function is not a component of the error, for, while the coefficients add to zero over every block, they do not do so over treatments (1CN) and (1CS). But if we take

$$z_1 - z_2 = (1CN)_1 - (1CS)_1 - (1CN)_2 + (1CS)_2$$

the conditions are satisfied. This expression shows how the difference between two treatments changes from one block to another. It is a measure of the effect of the blocks on the treatment difference, and is usually called an interaction of treatments with blocks. Such interactions are the typical components of error.

If z_i is the difference between (1CN) and (1CS) in the i th block, any comparison among the z_i satisfies the rules required for a component of the error. Hence the sum of squares of deviations of the z_i , when divided by 2, contributes 3 d.f. to the error s.s. A similar result holds if we take *any* comparison among the treatments and calculate it separately for each block. Since the 9 treatments provide 8 independent comparisons, we see that the 24 d.f. for the interactions of treatments with blocks may be divided into 8 sets of 3 d.f. The sum of squares for these 24 d.f. can of course be found by subtraction as shown in table 3.7.

TABLE 3.7 SUBDIVISION OF THE SUM OF SQUARES FOR ERROR

	d.f.	s.s.	m.s.
Error	36	544,690	15,130
Among controls	12	307,312	25,609
Treatments \times blocks	24	237,378	9,891

The F -ratio of the two components of error is 2.59, with 12 and 24 d.f. In the ordinary table this corresponds to a significance level of about 21½%. It is probably more correct to make a two-tailed test by doubling this probability. Either test indicates that the controls are more variable than the other treatments. This is not very surprising, since in data of this type the variance may tend to increase as the mean increases. In

any event the result throws doubt on the suitability of the original mathematical model; it would seem better to ascribe different error variances to the control and the fumigants.

The control also contributes to the treatments \times blocks component of the error. The 8 comparisons among the 9 treatments can be divided into 1 between the control and all fumigants, and 7 which are comparisons among the fumigants themselves. The former will contribute 3 d.f. to the treatments \times blocks interaction; the latter, 21 d.f. If the control is more variable than the fumigants, we would expect the mean square for the 3 d.f. to be larger than that for the 21 d.f., though it should be remembered that a mean square with only 3 d.f. is poorly determined. In the next section these two components will be computed separately.

3.63 Calculation of a Separate Error for a Treatment Comparison. As the eelworm example illustrates, certain treatments may be erratic in their effects, while others in the same experiment show more stable results. When this occurs, comparisons involving the erratic treatments have a higher experimental error variance than those among the more stable treatments. Since the error m.s. in the analysis of variance is some weighted average of these different variances, its use for individual t -tests may not be valid. In such cases it is helpful to be able to calculate a separate error for any specific treatment comparison. The procedure will be illustrated for the comparison between the control and the fumigants.

The first step is to compute the comparison separately for each block.

	Totals in block				
	1	2	3	4	Total
Control	1814	1295	1769	980	5858
Fumigants	2569	1780	2083	1490	8822
2 (Control)—(Fumigants)	1059	810	555	470	2894

Since there are 4 control plots and 8 fumigated plots in a block, the control total is multiplied by 2 in forming the comparison. The contribution to the error s.s. is

$$\frac{(1059)^2 + (810)^2 + (555)^2 + (470)^2}{24} - \frac{(2894)^2}{96} = 8862$$

The divisor 24 is found by the usual rule. The mean square is 2954, with 3 d.f. By subtraction, the mean square for the interaction of fumigants with blocks (21 d.f.) is found to be 10,882. Contrary to an-

ticipation, the component involving the control has the smaller mean square, though the difference does not approach significance.

As often happens when the simple model is inadequate, the best approach for a more accurate analysis is questionable. It is possible to compute a separate error, by the method just given, for each t -test that is made. This is the procedure least open to criticism, since it makes no assumption that errors are constant from one treatment to another. Since, however, each t -test would be based on only 3 d.f., these tests are insensitive, so that this method should not be used without good reason. In the present case it is probably justifiable to postulate only two error variances. The first, appropriate to the controls, is estimated by the mean square 25,609, with 12 d.f.; the second, for the fumigants, by the mean square 10,882, with 21 d.f. By this approach the variance of the mean response to the double dressing, for example, is estimated as

$$\frac{10,882}{16} + \frac{25,609}{16} = 2281$$

since each mean is taken over 16 plots.

Separation of the error s.s. into *single* components is seldom required except in special studies or as an exercise. One method is to set up 8 orthogonal comparisons among the treatments. Calculate each comparison separately in each block, and let z_{ij} be the value obtained for the i th comparison ($i = 1, 2, \dots, 8$) in the j th block. Then find

$$\begin{aligned}w_{i1} &= z_{i1} - z_{i2}; & w_{i2} &= z_{i1} + z_{i2} - 2z_{i3}; \\w_{i3} &= z_{i1} + z_{i2} + z_{i3} - 3z_{i4}\end{aligned}$$

Each w is a component of the error, and the whole 24 w 's are mutually orthogonal. The squares of the w 's, each with its proper divisor, give a separation of the error s.s. into single components.

3.7 Missing Data

3.71 Method of Handling Missing Data in the Analysis. From time to time certain observations are missing, through failure to record, gross errors in recording, or accidents. The omissions naturally affect the method of analysis. With each of the common designs we give computational instructions for analyzing data that contain gaps. The object of this section is to indicate the theoretical basis for these methods.

When certain observations are absent, the correct procedure is to write down a mathematical model for *all* observations that are present. The least squares normal equations are then constructed in the usual

way. These take exactly the same general form as when all observations are present; i.e.,

$$\sum (y_{ijk} - m - t_i - b_j) = 0 \quad (3.20)$$

over all observations whose equations contain any specified parameter that is to be estimated. Since, however, the terms in the equation corresponding to missing observations are absent, the system of equations loses some of the symmetry that it possesses when all observations are present, and the solutions are more difficult. The same general procedure supplies F - and t -tests of hypotheses about the nature of the treatment effects as described in section 3.24, though again the details become more complicated.

In the analysis of variance, two changes may be noted. Owing to the missing observations, the treatments and blocks s.s. become entangled, so that the treatments s.s. must be computed after allowing for block effects, as mentioned at the end of section 3.26. Secondly, if a observations are absent, the total number of degrees of freedom is reduced by a . Unless one or more complete treatments or blocks is missing, the number of parameters required to describe these effects will be the same as before. Consequently, the missing degrees of freedom all come from the error s.s., which now represents $(n_e - a)$ degrees of freedom. In short, missing data may be handled by applying the standard least squares procedure to all observations that are *not* missing. For future reference, this method will be called the "correct least squares procedure."

To the experimenter it may be a difficult business to carry out the construction and solution of a set of unfamiliar normal equations, even though he is quite competent to analyze a set of complete data. For this reason Yates (3.9), following a suggestion by Fisher, considered inserting values for the missing observations so as to obtain a set of complete data. Suppose that only a single observation is missing, and that a value x is substituted for this observation. If the analysis of variance is calculated in the usual manner for complete data, the error s.s. is found to be of the form

$$Ax^2 - 2Bx + C$$

where A , B , and C are numbers that are determined by the type of design and the values for the other observations (A is always positive).

In order to find a numerical value for x , Yates proposed to use the value that minimizes the error s.s. This is $x = B/A$. If this value is inserted in place of the missing observation, and if the data are analyzed as if no observations were absent, Yates showed that several important properties hold. (i) The estimates of treatment and block effects are

exactly the same as those obtained by the correct least squares procedure. (ii) The error s.s. is exactly the same as given by the correct procedure. (iii) To obtain the correct partition of the degrees of freedom, we subtract 1 from the total s.s. and 1 from the error s.s.

Yates also showed that the method of insertion fails to agree with the correct least squares procedure in two respects. The treatments s.s., as obtained in the analysis of variance of the "complete" data, is always slightly larger than the correct treatments s.s. for an F -test of the treatments. Unless an appreciable fraction of the total observations is missing, this overestimation is unlikely to be large; further, the exact F -test can be obtained by means of some additional calculations. The second defect of the method of insertion is that it may not give proper t -tests. That this will happen is clear, because in the analysis of "complete" data r replications are ascribed to the treatment that contains the missing observation, whereas there are only $(r - 1)$ replications. To allow for this disturbance we give special rules which provide t -tests that are approximately correct.

Thus the method adopted with a missing value is first to "estimate" this value by means of the formula B/A , which will be presented for the common designs. This estimate is used in place of the missing value, and the rest of the analysis is conducted as if the data were complete (except for the changes in degrees of freedom). Special methods are available for exact F -tests and for t -tests. If several observations are absent, a repeated application of the formula enables values to be substituted for each missing observation.

This method is essentially an ingenious computational device whose purpose is to enable the easy computations that apply to complete data to be used even when data are incomplete. Substitution of estimates for the missing data does not in any way recover the information that is lost through loss of data, as some experimenters have suggested, usually facetiously; it merely attempts to reproduce the results obtained by an application of the least squares method to the data that are present. The only complete solution of the "missing data" problem is not to have them.

3.8 The Analysis of Covariance

3.81 Purpose of the Technique. As indicated in section 3.2, experiments can be planned so that certain types of environmental effect are eliminated from the estimates of the treatment effects, with the result that these estimates are made more accurate. In the eelworm experiment, where the plots were grouped into blocks of 12 plots each, any dif-

ferences from block to block in the severity of eelworm infestation were eliminated in this way. On the other hand, differences in infestation from plot to plot within the same block are not controlled by the design and do contribute to the experimental errors, since treatments were assigned at random to plots within each block. Accordingly, before the fumigants were applied, samples were taken in order to estimate the natural infestation on each plot.

The analysis of covariance shows how to use these supplementary data to reduce the experimental errors by eliminating the effects of variations in the initial infestation within a block. The technique is potentially very useful. It often happens that some source of variation which cannot be controlled by the design can be measured by taking additional observations. Whenever this is so, the analysis of covariance can be utilized, often to great advantage. One caution is that, since the additional observations are to measure *environmental* effects, they must not be influenced by the treatments. The situation where such measurements are influenced by the treatments is discussed in section 3.88.

3.82 Initial Steps in the Analysis. The first step is to construct a new mathematical model. If y_{ijk} refers to the final eelworm count and x_{ijk} to the initial, the relation is

$$y_{ijk} = \mu + \tau_i + \beta_j + \gamma(x_{ijk} - \bar{x}) + e_{ijk} \quad (3.21)$$

The only change is the introduction of a new term to describe the effect of the initial eelworm number. We have assumed that the effect is linear, i.e., it is a constant multiple γ of the amount by which the initial eelworm number x_{ijk} on the plot differs from the average initial number \bar{x} for the whole experiment.

As before, the unknowns are estimated by least squares. In this case we minimize

$$\sum [y_{ijk} - m - t_i - b_j - c(x_{ijk} - \bar{x})]^2 \quad (3.22)$$

The normal equation for m is again of the form

$$\sum [y_{ijk} - m - t_i - b_j - c(x_{ijk} - \bar{x})] = 0 \quad (3.23)$$

over all observations. If the same linear restrictions as before are applied to the t_i and the b_j , equation (3.23) implies that m is the mean of the y_{ijk} . The equation for t_i is the same as (3.23) except that the sum is over those observations that receive t_i . The equation may be rearranged as

$$r_{im} + r_{it_i} = T_{iy} - c(T_{ix} - r_{i\bar{x}}) \quad (3.24)$$

where r_i is the number of replications, and T denotes a treatment total. This gives

$$m + t_i = \frac{T_{iy}}{r_i} - c \left(\frac{T_{ix}}{r_i} - \bar{x} \right) \quad (3.25)$$

The important feature of this equation is that in order to obtain t_i , the observed treatment mean of the y 's (T_{iy}/r_i) is *adjusted*, the adjustment depending on the treatment mean of the x 's. It is this adjustment that removes the effect of the initial infestation.

The equation for c , the adjustment factor, will not be developed in full. It turns out that c is given by the ratio of the error sum of products of y and x to the error s.s. of x .

3.83 Computations. In practice we start with a joint analysis of the sums of squares and products of y and x . In order to illustrate certain features of the covariance technique, the treatments s.s. will be subdivided into "linear" and "curvature" components, as was done for y in table 3.5 (p. 64). The original values for x are given above the y values in table 3.1 (p. 40). The treatment totals and the quantities

TABLE 3.8 TREATMENT TOTALS FOR NUMBERS OF CYSTS *

Level	Before fumigation (x)					After fumigation (y)				
	CN	CS	CM	CK	Total	CN	CS	CM	CK	Total
0			1975		1975			5858		5858
1	402	417	513	570	1902	1066	928	1431	892	4317
2	389	554	568	778	2289	1265	877	1241	1122	4505
2(2) + (1):L	1180	1525	1649	2126	6480	3596	2682	3913	3136	13327
2(1) - (2):Q	415	280	458	362	1515	867	979	1621	662	4129

* Numbers of cysts per plot were the numbers found in 400 grams of soil. Thus the totals in the table represent 1600 grams of soil, except for the no-fumigant totals, which represent 6400 grams.

needed for isolating the linear response and the curvature are reproduced in table 3.8. The analysis of the sum of squares of x follows by the same methods as given previously for y .

To analyze the sum of products, we carry out the same operations as for a sum of squares, except that at every stage a square is replaced by the corresponding product. A few examples will suffice.

$$\text{Total: } (269)(466) + (283)(280) + \dots + (48)(298) - \frac{(6166)(14,680)}{48} = 355,929$$

$$\text{Treatments: } \frac{(1975)(5858)}{16} + \frac{(402)(1066)}{4} + \dots + \frac{(778)(1122)}{4} - \frac{(6166)(14,680)}{48} = -9222$$

$$\text{Average linear response: } \frac{(2289 - 1975)(4505 - 5858)}{32} = -13,276$$

The complete analysis of covariance is shown in table 3.9.

TABLE 3.9 SUMS OF SQUARES AND PRODUCTS (x = BEFORE, y = AFTER, FUMIGATION)

	d.f.	(x^2)	(yx)	(y^2)
Blocks	3	159,618	175,873	289,427
Treatments	8	29,142	-9,222	157,448
Average linear response	1	3,081	-13,276	57,207
Average curvature	1	2,204	8,285	31,140
Differences in linear	3	22,975	-6,837	43,408
Differences in curvature	3	882	2,606	25,693
Error	36	121,408	189,278	544,690
Total	47	310,168	355,929	991,565

If we denote the sums of squares and products for error by E_{xx} , E_{yy} , and E_{yx} , respectively, the adjustment factor c , or regression coefficient of y on x , is given by

$$c = \frac{E_{yx}}{E_{xx}} = \frac{189,278}{121,408} = 1.559024$$

It may seem surprising that the coefficient is greater than 1. This is probably explained by the great seasonal increase in the eelworm numbers, obvious from table 3.8.

The residual error s.s. may now be found. The original error s.s. is 544,690, with 36 d.f. To remove the effect of the regression on the initial eelworm numbers, we subtract

$$\frac{E_{yx}^2}{E_{xx}} = \frac{(189,278)^2}{121,408} = 295,089$$

$$E_{yy} - \frac{E_{yx}^2}{E_{xx}} = 544,690 - 295,089 = 249,601$$

Thus the residual s.s. is 249,601, with 35 d.f., since 1 d.f. must be sub-

where r_i is the number of replications, and T denotes a treatment total. This gives

$$m + t_i = \frac{T_{iy}}{r_i} - c \left(\frac{T_{ix}}{r_i} - \bar{x} \right) \quad (3.25)$$

The important feature of this equation is that in order to obtain t_i , the observed treatment mean of the y 's (T_{iy}/r_i) is *adjusted*, the adjustment depending on the treatment mean of the x 's. It is this adjustment that removes the effect of the initial infestation.

The equation for c , the adjustment factor, will not be developed in full. It turns out that c is given by the ratio of the error sum of products of y and x to the error s.s. of x .

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2(2) + (1):L	1180	1525	1649	2126	6480	3596	2682	3913	3136	13327
2(1) - (2):Q	415	280	458	362	1515	867	979	1621	662	4129

* Numbers of cysts per plot were the numbers found in 400 grams of soil. Thus the totals in the table represent 1600 grams of soil, except for the no-fumigant totals, which represent 6400 grams.

needed for isolating the linear response and the curvature are reproduced in table 3.8. The analysis of the sum of squares of x follows by the same methods as given previously for y .

To analyze the sum of products, we carry out the same operations as for a sum of squares, except that at every stage a square is replaced by the corresponding product. A few examples will suffice.

$$\text{Total: } (269)(466) + (283)(280) + \dots + (48)(298) - \frac{(6166)(14,680)}{48} = 355,929$$

$$\text{Treatments: } \frac{(1975)(5858)}{16} + \frac{(402)(1066)}{4} + \dots + \frac{(778)(1122)}{4} - \frac{(6166)(14,680)}{48} = -9222$$

$$\text{Average linear response: } \frac{(2289 - 1975)(4505 - 5858)}{32} = -13,276$$

The complete analysis of covariance is shown in table 3.9.

TABLE 3.9 SUMS OF SQUARES AND PRODUCTS (x = BEFORE, y = AFTER, FUMIGATION)

	d.f.	(x^2)	(yx)	(y^2)
Blocks	3	159,618	175,873	289,427
Treatments	8	29,142	-9,222	157,448
Average linear response	1	3,081	-13,276	57,207
Average curvature	1	2,204	8,285	31,140
Differences in linear	3	22,975	-6,837	43,408
Differences in curvature	3	882	2,606	25,693
Error	36	121,408	189,278	544,690
Total	47	310,168	355,929	991,565

If we denote the sums of squares and products for error by E_{xx} , E_{yy} , and E_{yx} , respectively, the adjustment factor c , or regression coefficient of y on x , is given by

$$c = \frac{E_{yx}}{E_{xx}} = \frac{189,278}{121,408} = 1.559024$$

It may seem surprising that the coefficient is greater than 1. This is probably explained by the great seasonal increase in the eelworm numbers, obvious from table 3.8.

The residual error s.s. may now be found. The original error s.s. is 544,690, with 36 d.f. To remove the effect of the regression on the initial eelworm numbers, we subtract

$$\frac{E_{yx}^2}{E_{xx}} = \frac{(189,278)^2}{121,408} = 295,089$$

$$E_{yy} - \frac{E_{yx}^2}{E_{xx}} = 544,690 - 295,089 = 249,601$$

Thus the residual s.s. is 249,601, with 35 d.f., since 1 d.f. must be sub-

tracted for the additional parameter γ . The residual m.s. s_{yx}^2 is 249,601/35, or 7131. This is less than half the original mean square of 15,130, and indicates that the use of covariance has approximately doubled the accuracy of the experiment.

One feature of the covariance method deserves comment at this point. In the model the effect of x was assumed to be linear, but no assumption was made about the *strength* of the effect. That is left to be determined by the data. If the eelworm numbers on each plot had remained unchanged throughout the season, except for the influence of the treatments, and if the eelworm numbers had been accurately measured, the residual s.s. would have been zero. The residual s.s. as found presumably represents a contribution due to seasonal variations in infestation and one due to the fact that the eelworm numbers were not estimated accurately, being obtained only from small samples of soil. If the variable x had had *no* linear effect on y , the residual m.s. would have been the same as the original mean square, apart from sampling fluctuations.

3.84 The Adjusted Treatment Means. A little time is saved if the adjusted treatment totals are found first. From equation (3.24) the adjustment to the i th treatment total of y is

$$-c(T_{ix} - r_i\bar{x})$$

where r_i is the number of replicates, and \bar{x} the general mean of x (128.46). Thus for the control the adjusted total is

$$5858 - 1.5590(1975 - 16 \times 128.46) = 5858 + 125 = 5983$$

and for (1CN)

$$1066 - 1.5590(402 - 4 \times 128.46) = 1066 + 174 = 1240$$

The *means* are shown in table 3.10.

TABLE 3.10 ADJUSTED TREATMENT MEANS (EELWORMS PER 400 GM. SOIL)

Level of dressing	Fumigant			
	CN	CS	CM	CK
0			374	
1	310	270	358	201
2	365	204	289	178

3.85 t -tests. These tests are slightly complicated by the fact that account must be taken of the sampling error of the adjustment factor c . The difference between two adjusted means may be written

$$\bar{y}_1 - \bar{y}_2 - c(\bar{x}_1 - \bar{x}_2)$$

From regression theory the variance of this quantity is given by

$$\sigma_{y \cdot x}^2 \left(\frac{1}{r_1} + \frac{1}{r_2} \right) + (\bar{x}_1 - \bar{x}_2)^2 \sigma_c^2 = \sigma_{y \cdot x}^2 \left[\frac{1}{r_1} + \frac{1}{r_2} + \frac{(\bar{x}_1 - \bar{x}_2)^2}{E_{xx}} \right] \quad (3.26)$$

where $\sigma_{y \cdot x}^2$ is the residual error variance, and E_{xx} the error s.s. for x . As an estimate of $\sigma_{y \cdot x}^2$ we use the residual error m.s., in this case 7131 with 35 d.f.

Example. A t -test of the reduction to the double dressing of CS. The reduction is $(374 - 204)$, or 170 eelworms per sample. The estimated variance is

$$(7131) \left[\frac{1}{16} + \frac{1}{4} + \frac{(123.44 - 138.50)^2}{121,408} \right] = (7131)(0.31437) = 2242 \quad (3.27)$$

the means of the x 's being found by division from table 3.8. Hence

$$t = \frac{170}{\sqrt{2242}} = \frac{170}{47.35} = 3.59$$

with 35 d.f., which is highly significant.

For a more complicated comparison of the adjusted means,

$$\sum l_i [\bar{y}_i - c(\bar{x}_i - \bar{x})]$$

the estimated variance is

$$s_{y \cdot x}^2 \left[\sum \frac{l_i^2}{r_i} + \frac{(\sum l_i \bar{x}_i)^2}{E_{xx}} \right] \quad (3.28)$$

(The term in \bar{x} vanishes because for any comparison the sum of the l 's must be zero.)

One annoying feature of these tests is that the x values enter into the variance, so that every comparison necessitates a separate computation of the variance. As a time-saving approximation, Finney (3.10) has proposed that an average value for the contribution of the term in the x 's may be used. This amounts to using

$$s_{y \cdot x}^2 \left[1 + \frac{t_{xx}}{E_{xx}} \right] = (7131) \left[1 + \frac{3643}{121,408} \right] = (7131)(1.030) = 7345$$

as the *effective* residual error m.s., where t_{xx} is the treatments *mean square* for x . The term in brackets represents the average contribution from the x 's, or in other words the contribution from sampling errors in the adjustment factor c .

Thus, for the variance of the difference between the adjusted means of the control and 2CS, we would use

$$(7345) \left(\frac{1}{r_1} + \frac{1}{r_2} \right) = (7345) \left(\frac{1}{16} + \frac{1}{4} \right) = 2295$$

instead of the value 2242 given by the more exact expression (3.27). Similarly, instead of (3.28), we use

$$(7345) \sum \left(\frac{l_i^2}{r_i} \right)$$

This approximation is usually good enough if the number of error degrees of freedom exceeds 20, since in such cases the contribution from errors in c is small. The more exact test may be used if n_e is small.

3.86 F-tests. The fact that the same adjustment factor appears in every treatment mean also influences F -tests. The procedure is shown in table 3.11.

TABLE 3.11 F -TEST WITH THE ANALYSIS OF COVARIANCE

	d.f.	(x^2)	(ηx)	(η^2)	Residuals		
					d.f.	s.s.	m.s.
Treatments	8	29,142	-9,222	157,448	8	237,192	29,649
Error	36	121,408	189,278	544,690	35	249,601	7,131
T + E	44	150,550	180,056	702,138	43	486,793	

The figures to the left of the vertical line are from the previous analysis (table 3.9), and in practice would not be recopied. Form a new line in the analysis by addition of the items for treatments and error. From this value for (y^2) , i.e., 702,138, subtract the contribution due to a regression on x ,

$$\frac{(180,056)^2}{150,550} = 215,345$$

The remainder, 486,793, is entered in the column headed "residuals s.s." and carries 43 d.f., 1 being subtracted for the regression. The same process is completed for the error line; actually, this was already done in computing the residual error s.s. The residual s.s. for treatments is found by subtracting that for error from that for treatments + error

$$237,192 = 486,793 - 249,601$$

It always has the same number of degrees of freedom as the original treatments s.s. The F -test of the adjusted treatment means is given by the ratio of the residual m.s.

$$F = \frac{29,649}{7131} = 4.16$$

with 8 and 35 d.f.

If it is desired to test some component of the treatments s.s., the same calculation is made with the component in place of the treatments s.s. throughout. If several components are to be tested, this becomes rather tedious. A useful approximation is to construct from the original analysis of covariance in table 3.9 an analysis of $(y - cx)^2$. This is most easily done by multiplying each term in (x^2) by c^2 , or 2.43048, each term in (yx) by $-2c$, or -3.11805 , and adding the two products to (y^2) . The results are shown in table 3.12.

TABLE 3.12 ANALYSIS OF VARIANCE OF $(y - cx)$

	d.f.	s.s.	m.s.	F	F'
Treatments	8	257,032	32,129		
Average linear response	1	106,090	106,090	14.88	14.51
Average curvature	1	10,664	10,664	1.50	1.47
Differences in linear	3	120,566	40,189	5.64	5.05
Differences in curvature	3	19,711	6,570	0.92	0.92
Error	35	249,592	7,131		

It will be noted that this calculation gives the correct residual error m.s. However, sums of squares for components of the treatments are always larger than those given by the more roundabout correct procedure. Thus the F values shown on the right are all too large. The overestimation is seldom great, and the F values serve for a preliminary inspection. Those that are just beyond the significance level may be recomputed by the correct procedure if it is thought worth while. For comparison, the F' values shown above are those obtained by the correct procedure (it is a useful exercise to check them). With either F or F' there is a significant average linear response and significant differences among fumigants in their linear responses, while the curvature terms do not approach significance. In the analysis made without covariance (table 3.5, p. 64) no component was significant.

3.87 The Increase in Accuracy Due to Covariance. From table 3.9 we see that with no covariance the error s.s. for y is 544,690, with 36 d.f., giving an error m.s. of 15,130. A comparable figure when covariance is used is the effective residual m.s., 7345. This is preferable to the

residual m.s. itself (7131) because it makes allowance for the sampling error of the adjustment factor c . The accuracy obtained with covariance relative to that without covariance is estimated by $15,130/7345$, or 2.06. The use of covariance appears to have had about the same effect as doubling the number of replicates. In making this comparison, we ignored the effect of the reduction in error d.f. from 36 to 35, because it is negligible.

As in ordinary regressions, the x variable may be transformed to another scale if this is likely to produce a more linear relation with y . The use of $\log x$, for instance, is common in biological work. Two or more different x variables may be used. The calculations for this case are described by Snedecor (3.4, chapter 13).

3.88 The Case Where the x Variable is Influenced by the Treatments.

In a covariance analysis the treatment mean \bar{y}_i is adjusted by the amount $-c(\bar{x}_i - \bar{x})$. The effect of the adjustment is to change each \bar{y}_i to the value that it would be expected to have if all treatments had the same x mean. It is in this way that the technique removes the effect of variations in the \bar{x}_i . If, however, the differences among the \bar{x}_i are in part produced by the treatments, the adjustment removes part of the treatment effect and its interpretation is changed. Consequently, in the standard use of covariance, it is important to be sure that the treatments did not affect the x values. Sometimes this is obvious, as in the eelworm experiment, because the x values were recorded before the treatments were applied. A more doubtful case is that of the number of plants per plot in a field experiment, counted after the application of the treatments, which may or may not influence plant numbers. An F -test of the x values is helpful in such cases.

Where the treatments do affect x , a covariance analysis may add information about the way in which the treatments produced their effects. In the eelworm experiment, the yields of the oats which were grown on the plots were obtained. Since eelworms attack oats, it would be interesting to know whether the effects of the treatments on the oats were simply a reflection of their effects on the eelworms. This is examined by a covariance analysis in which the oats yields are the y values, and the eelworm numbers after harvest are the x values. If the F -test of the adjusted treatment means still shows significance, the conclusion is that not all the treatment effect on the oats can be attributed to the reduction in eelworm numbers. This happened in the present instance, because some treatments supplied nitrogen to the crop, and therefore acted in part as fertilizers as well as fumigants. As Bartlett (3.11) has pointed

out, the interpretation of this use of covariance requires care, since a hidden extrapolation may be involved.

When the x variables are influenced by treatments, the short-cut method for t -tests by use of the effective residual error and the approximate F -tests by means of an analysis of $(y - \bar{x})$ should be avoided, since they may be seriously in error.

3.9 Effects of Errors in the Assumptions Underlying the Analysis of Variance

The assumptions made in the analysis of variance are that treatment and environmental effects are additive, and that the experimental errors are independently distributed in the normal distribution, with a common variance. In practice we can never be sure that these assumptions all hold, and often there is good reason to suspect that some are false. The consequences of failures in the assumptions and the remedial steps to be taken have been summarized recently by Eisenhart (3.12), Cochran (3.13), and Bartlett (3.14). Only a few comments will be given here.

As a rule, the failure of an assumption will affect both the significance levels and the sensitivity of F - and t -tests. When the experimenter thinks that he is testing at the 5% level, he may actually be testing at the 8% level. Usually, though not invariably, the true significance probability is larger than the apparent one; that is, too many significant results are obtained. Also, there is usually a loss of sensitivity, in the sense that a more powerful test than the analysis of variance F -test could be constructed if the correct mathematical model were known. There is a corresponding loss of accuracy in the estimates obtained for the treatment effects, since these, too, could be made more accurate if the correct model were known.

Although generalization is hazardous, experience suggests that in the majority of experiments, at least in the field of biology, these disturbances are not sufficiently great to invalidate the technique. They do imply, however, that significance levels and confidence limits must be considered approximate rather than exact. For the same reason, the inflexible use of say the 5% significance level to divide the effects into those that are regarded as "real" and those that are not is hardly justifiable.

The most serious disturbances appear to arise when the experimental error variance is not constant over all observations. Sometimes this happens, as mentioned previously, because certain treatments are erratic in their effects. In such cases, the appropriate error variance for com-

paring one pair of treatments might be four times as large as that for another pair, and the use of the same estimated variance for both comparisons would lead to t -tests that were completely erroneous. Where this type of disturbance is suspected, the remedy is to divide the error s.s. into components each of which is homogeneous, as indicated in section 3.63.

The same problem may arise because the experimental errors follow a distribution that is decidedly skew. In such distributions the error variance for a treatment tends to be a function of the mean produced by the treatment. If the nature of the functional relationship is known, a transformation can be found that will place the data on a scale on which the error variance is more nearly constant. This transformation is then made on the observations before starting the analysis. The principal transformations that have been found useful are discussed by Bartlett (3.14); they include logs, square roots, and (for data expressed in fractions) inverse sines.

Such transformations may also be useful in cases where treatment and environmental effects are not additive. If, for instance, a treatment increases all observations by 20%, irrespective of the initial level, a change to logs will introduce additivity. When transformations are made for this purpose, it should be realized that they will also affect the distribution of the experimental errors. Fortunately, it often happens that such transformations also bring the distribution of errors closer to normality.

Finally, the assumption that the errors are independent from observation to observation may be obviously untenable. It is well known that crop yields on neighboring plots tend to be positively correlated, and in laboratory experiments observations made by the same person at about the same time tend to exhibit the same type of correlation. These correlations might completely vitiate tests of significance. The remedy in this case is the proper use of randomization, which, as it were, introduces independence in the assignment of treatments to the experimental units or in the assignment of the order in which observations are made, so that the errors may effectively be regarded as independent. For further discussion of this question, see Yates (3.15), Fisher (3.3), Bartlett (3.16), and Welch (3.17).

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CHAPTER 4

COMPLETELY RANDOMIZED, RANDOMIZED BLOCK, AND LATIN SQUARE DESIGNS

4.1 Completely Randomized Designs

4.11 Description. The simplest type of layout is that in which treatments are allotted to the units entirely by chance. More specifically, if a treatment is to be applied to four units, for example, the randomization gives every group of four units in the experimental material an equal probability of receiving the treatment. In addition the units should be processed in random order at all subsequent stages of the experiment where this order is likely to affect the results.

This design has several conveniences:

1. Complete flexibility is allowed. Any number of treatments and of replicates may be used. The number of replications can be varied at will from treatment to treatment (though such variation is not recommended without good reason). All the available experimental material can be utilized—an advantage in small preliminary experiments where the supply of material is scarce.

2. The statistical analysis is easy even if the numbers of replicates are not the same for all treatments or if the experimental errors differ from treatment to treatment.

3. The method of analysis remains simple when the results from some units or from whole treatments are missing or are rejected. Moreover, the relative loss of information due to missing data is smaller than with any other design.

The principal objection to a completely randomized design is on the grounds of accuracy. Since the randomization is not restricted in any way to ensure that the units which receive one treatment are similar to those which receive another treatment, the whole of the variation among the units enters into the experimental error. For this reason the error can often be reduced by the use of a different design, unless the units are highly homogeneous or the experimenter has no information by which to arrange or handle the units in more homogeneous groups.

Complete randomization seems the obvious procedure for many lab-

oratory experiments, e.g., in physics, chemistry, or cookery, where a quantity of material, after thorough mixing, is divided into small samples or batches to which the treatments are applied. On the other hand, these designs are seldom used in field experiments, the method of randomized blocks having been found consistently more accurate.

One fact compensates to some extent for the higher experimental errors as compared with other designs. For a given number of treatments and a given number of experimental units, complete randomization provides the maximum number of degrees of freedom for the estimation of error. As pointed out in section 2.31, the sensitivity of the experiment increases as the number of error degrees of freedom is increased. Consider, for example, an experiment with 2 treatments in 4 replicates. The degrees of freedom for error are 6 under complete randomization as against 3 if the method of pairing (section 4.2) is used. From section 2.31 it appears that the paired experiment must produce about a 14% reduction in the error variance in order to offset the additional unreliability in the estimate of error. This point is worth bearing in mind with small experiments.

To summarize, complete randomization may be appropriate (i) where the experimental material is homogeneous, (ii) where an appreciable fraction of the units is likely to be destroyed or to fail to respond, and (iii) in small experiments where the increased accuracy from alternative designs does not outweigh the loss of error degrees of freedom.

4.12 Randomization. If the number of units does not exceed 16, tables 15.6 and 15.7 may be used. Suppose that there are 3 treatments, of which two have 4 replicates while the third has 8 replicates. Numbers 1-16 are assigned to the units in any convenient order. A random permutation is drawn from table 15.7, say

9, 13, 8, 5, 12, 1, 14, 16, 6, 7, 3, 4, 10, 11, 15, 2

The first treatment is applied to units 9, 13, 8, 5; the second to units 12, 1, 14, 16; and the third to the remainder.

With more than 16 units in the experiment, numbered discs, beans, or a book of random numbers may be used, as described in chapter 15.

4.13 Statistical Analysis. Table 4.1 shows some of the results of an experiment on the effects of applications of sulphur in reducing scab disease of potatoes. The object in applying sulphur is to increase the acidity of the soil, since scab does not thrive in very acid soil. In addition to untreated plots which serve as a control, 3 amounts of dressing were compared—300, 600, and 1200 lb. per acre. Both a fall and a spring ap-

plication of each amount was tested, so that in all there were 7 distinct treatments. The sulphur was spread by hand on the surface of the soil, and then disced in to a depth of about 4 inches. The quantity to be

TABLE 4.1 FIELD PLAN AND SCAB INDICES FOR A COMPLETELY RANDOMIZED EXPERIMENT ON POTATOES

F3 9	0 12	S6 18	F12 10	S6 24	S12 17	S3 30	F6 16
0 10	S3 7	F12 4	F6 10	S3 21	0 24	0 29	S6 12
F3 9	S12 7	F6 18	0 30	F6 18	S12 16	F3 16	F12 4
S3 9	0 18	S12 17	S6 19	0 32	F12 5	0 26	F3 4

Notation: F = fall, S = spring application, 0 = control. The numbers 3, 6, 12 are the amounts of sulphur in 100 lb. per acre.

Results grouped by treatments

	0	F3	S3	F6	S6	F12	S12
	12 30	9	30	16	18	10	17
	10 18	9	7	10	24	4	7
	24 32	16	21	18	12	4	16
	29 26	4	9	18	19	5	17
Totals	181	38	67	62	73	23	57
Means	22.6	9.5	16.8	15.5	18.2	5.8	14.2

$G = 501$

Analysis of variance

Source of variation	d.f.	s.s.	m.s.	F
Treatments	$(t - 1) = 6$	972.3	162.0	3.61 *
Error	$(N - t) = 25$	1122.9	44.9	
Total	$(N - 1) = 31$	2095.2		

* Denotes significance at the 5% level.

analyzed is the "scab index." This is, roughly speaking, the percentage of the surface area of the potato that is infected with scab. It is obtained by examining 100 potatoes at random from each plot, grading each potato on a scale from 0 to 100% infected, and taking the average.

The design was a completely randomized one, with 4 replications of each sulphur dressing and 8 replications of the control, which received extra replication in order to obtain a fairly good estimate of the natural infestation. Incidentally, a randomized blocks layout might have been superior.

The computations are very simple. Let y denote an observation, T_i a treatment total, G the grand total, r_i the number of replications of the i th treatment, $N = \sum r_i$ the total number of observations, and t the number of treatments.

✓ *Step 1.* Find the treatment totals and the grand total.

✓ *Step 2.* The sums of squares are computed as follows.

$$\text{Correction factor: } C = \frac{G^2}{N} = \frac{(501)^2}{32} = 7843.8$$

$$\text{Total: } \sum y^2 - C = (9)^2 + (12)^2 + \dots + (4)^2 - 7843.8 = 2095.2$$

$$\text{Treatments: } \sum \frac{T_i^2}{r_i} - C = \frac{(181)^2}{8} + \frac{(38)^2 + (67)^2 + \dots + (57)^2}{4} - 7843.8 = 972.3$$

$$\text{Error: (Total s.s.)} - (\text{treatments s.s.}) = 2095.2 - 972.3 = 1122.9$$

The analysis of variance is given at the foot of table 4.1. The F -ratio for treatments is significant at the 5% level. From the treatment means it appears that all dressings had some beneficial effect, and that the fall application was more effective than the spring one. There is little or no evidence that the higher dressings were more effective than the lowest dressing. If the summary is conducted from this point of view, we should isolate and test two individual components of the treatments s.s.: (i) a component measuring the average effect of all dressings, (ii) a component comparing the fall and spring applications. The following computations are required.

Average effect of sulphur. The total over all dressings is 320, representing 24 plots. Since the control total, 181, represents 8 plots, the comparison is

$$3(181) - 320 = 223$$

The contribution to the sum of squares in the analysis of variance is $(223)^2/96$, or 518.0. By the rules in section 3.42, the divisor, 96, is $[(9)(8) + 24]$.

Fall versus spring application. The comparison is

$$(38 + 62 + 23 - 67 - 73 - 57) = -74$$

It is obviously orthogonal to the previous comparison. The contribution to the sums of squares is $(74)^2/24$, or 228.2. These calculations lead to the analysis of variance in table 4.2.

The remaining four components of the treatments s.s. must represent comparisons among the levels of sulphur. The average reduction in scab due to sulphur is significant at the 1% level, while the superiority of the fall application is also significant. Differences among the levels show no sign of significance.

TABLE 4.2 SUBDIVISION OF THE TREATMENTS SUM OF SQUARES IN THE EXPERIMENT ON POTATO SCAB

Source of variation	d.f.	s.s.	m.s.	F
Treatments	6	972.3	162.0	3.61 *
Control vs. sulphur	1	518.0	518.0	11.54 **
Fall vs. spring application	1	228.2	228.2	5.08 *
Comparisons among levels	4	226.1	56.5	1.26
Error	25	1122.9	44.9	

* Denotes significance at the 5% level.

** Denotes significance at the 1% level.

The conclusions might be phrased as follows. "The application of sulphur produced a significant decrease in the scab index, the averages being 22.6 for the untreated plots, 16.4 for plots with the spring application, and 10.2 for plots with the fall application. The fall application proved significantly better than the spring application. There was no indication that the higher levels of dressing were more effective than the lowest level." It should be remarked that a continuation of this experiment and other experiments caused these conclusions to be modified, the higher levels having shown to better effect.

A more extensive discussion of the analysis of experiments of this type is given by Snedecor (4.1). Note that, if certain observations are missing or have to be rejected, the method of computation remains exactly the same except for the slight complication that different treatments will usually have different numbers of replications.

The practice of calculating a separate standard error for each treatment is still found in some types of experimentation. Unless the treatments have different error variances, the use of a pooled error, as given by the analysis of variance, is recommended since more sensitive tests of significance are obtained. In cases where the experimental error appears to vary considerably from treatment to treatment, a test of homogeneity of the error variance, due to Bartlett (4.2), may be made. This test is also described in Snedecor (4.1, section 10.13).

It is worth emphasizing that, even if the treatments are randomly assigned to the units, the above methods of analysis do not apply when restrictions on the randomization are introduced at a later stage in the experiment. Consider a chemical experiment in which samples of liquid

from the same bottle are treated in four different ways. The samples, being mutually indistinguishable, are assigned at random to the treatments. It is, however, practicable to treat only a few samples during each work period, and the investigator decides to complete a single replication at each session. The result of this additional control is that the differences among replicates are eliminated from the experimental errors and must equally be eliminated from the estimated errors. The analysis should follow the method described in section 4.2. This point will arise frequently in experiments where many physical operations must be carried out.

4.14 Standard Errors. The estimated standard error of the difference between two treatment means is

$$s_d = \sqrt{s^2 \left(\frac{1}{r_1} + \frac{1}{r_2} \right)}$$

This formula applies when a pooled error is used, s^2 being the pooled error m.s. per unit, and r_1 and r_2 the numbers of replicates for the two treatments. If $r_1 = r_2 = r$, the formula reduces to $\sqrt{\frac{2s^2}{r}}$. The degrees of freedom for t -tests are those in s^2 .

Example. The average effect of the sulphur has already been tested by an F -test. We will perform the same test by means of a t -test applied to the treatment means. Since the control has 8 replications, and the mean of all sulphur dressings has 24 replications,

$$s_d = \sqrt{(44.9) \left(\frac{1}{8} + \frac{1}{24} \right)} = 2.736$$

The mean scab index for the control is 22.62, and that for the dressings is 13.33. Hence

$$t = \frac{22.62 - 13.33}{2.736} = 3.395$$

with 25 d.f. The value of t^2 is 11.53, in agreement with the F found in the analysis of variance.

If the true error variances per unit are considered to be different for the two treatments, the appropriate formula is

$$s_d = \sqrt{\frac{s_1^2}{r_1} + \frac{s_2^2}{r_2}}$$

where s_1^2 , s_2^2 are the respective error m.s. per unit for the two treatments.

In this case the ratio of the treatment difference to s_d does not follow Student's t -distribution except in special instances. The development of correct significance levels for this case presents a problem that has stimulated much discussion in recent years. Various methods that have been suggested agree closely in practical application except when the numbers of degrees of freedom in s_1^2 and s_2^2 are small. Fisher and Yates (4.3, tables V1 and V2) give tables of the significance levels of d/s_d derived from Fisher's theory of fiducial probability. (Note that their s_1^2 corresponds to our s_1^2/r_1 , etc.) Where these tables are not readily accessible, the following approximation is suggested. It probably errs slightly on the conservative side, in the sense that the value of t required for significance may be slightly too high.

Let n_1, n_2 be the numbers of degrees of freedom in s_1^2, s_2^2 , respectively. From the ordinary t -table record the significance levels t_1, t_2 corresponding to n_1 and n_2 degrees of freedom, respectively. The approximate significance level for the ratio d/s_d is

$$t' = \frac{w_1 t_1 + w_2 t_2}{w_1 + w_2}$$

where

$$w_1 = \frac{s_1^2}{r_1}; \quad w_2 = \frac{s_2^2}{r_2}$$

Since t' always lies between the ordinary t values for n_1 and n_2 degrees of freedom, this calculation is needed only for those occasional cases where d is close to the borderline of significance. Further, when $n_1 = n_2 = n$, t' is the ordinary t value for n degrees of freedom.

Example. This experiment is a case where one might suspect that the error variance would not be homogeneous. If a treatment is very successful in reducing scab, the scab indices for that treatment will all be close to zero; hence their variance must be small. With a treatment such as the control the scab index has a greater possible range of variation. Consequently, we might expect the error variance for a treatment to depend on the mean produced by the treatment. With only 3 d.f. for the estimated error variance per treatment, we cannot hope to detect small differences in the true error variances. Perhaps the most important comparison is that between the variance for the control and the average variance for the sulphur dressings. The error m.s. will be found to be 70.0 for the control (with 7 d.f.) and 35.2 for the dressings (with 6×3 or 18 d.f.). The F -ratio, 1.99, falls just short of the 10% level of significance.

Consequently, the conclusions will be on a sounder basis if we ascribe different true error variances to the control and the dressings. For a

t-test of the average effect of sulphur we now take

$$s_d = \sqrt{\frac{70.0}{8} + \frac{35.2}{24}} = \sqrt{8.75 + 1.47} = 3.197$$

The value of *t* is 9.29/3.197, or 2.906. To test this at the 1% level by the approximate test above we have

$$n_1 = 7; \quad n_2 = 18; \quad t_1 = 3.499; \quad t_2 = 2.878; \quad w_1 = 8.75; \quad w_2 = 1.47$$

Hence the value required for significance is

$$t' = \frac{(8.75)(3.499) + (1.47)(2.878)}{10.22} = 3.410$$

The average effect of sulphur now falls short of the 1% significance level. It remains significant at the 5% level. For comparisons among the different sulphur dressings we would use ordinary *t*-tests, where *s*² is taken as 35.2 with 18 d.f. The effect will be to enhance the significance of comparisons among the dressings, though the conclusions quoted previously are not altered.

As mentioned in section 3.9, it is usually better to handle this type of experiment, where the error variance is thought to be a function of the mean, by converting the original data to a scale on which the error vari-

TABLE 4.3 ANALYSIS OF VARIANCE OF THE SQUARE ROOTS OF THE SCAB INDICES

Source of variation	d.f.	s.s.	m.s.	<i>F</i>
Treatments	6	18.22	3.04	4.19
Control vs. sulphur	1	8.17	8.17	11.27
Fall vs. spring application	1	4.59	4.59	6.33
Comparisons among levels	4	5.46	1.36	1.88
Error	25	18.12	0.725	
Error for controls	7	6.27	0.896	
Error for dressings	18	11.85	0.658	
Total	31	36.34		

ances will be more nearly constant. On the supposition that the error variance may be proportional to the mean, the square roots of the scab indices might have been analyzed. As a matter of interest this analysis is shown in table 4.3; square roots were recorded to only one decimal place.

The error m.s. for the controls is now closer to that for the dressings, the *F*-ratio being 1.36 against 1.99 in the original analysis. Further, the *F*-ratios for most components of the treatments s.s. are higher than in the original analysis, suggesting an increase in the sensitivity of the analysis.

4.2 Single Grouping: Randomized Blocks

Each group
constitutes
a design
in all rep.

4.21 Description. The essence of this design is that the experimental material is divided into groups, each of which constitutes a single trial or replication. At all stages of the experiment the object is to keep the experimental errors within each group as small as is practicable. Thus, when the units are assigned to the successive groups, all units which go in the same group should be closely comparable. Similarly, during the course of the experiment, a uniform technique should be employed for all units in the same group. Any changes in technique or in other conditions that may affect the results should be made between groups.

This division into replications need be recognized only at those stages in the conduct of the experiment where the division may help to reduce experimental errors. In agricultural field experiments the division is made at the start when the plots are marked out in the field. Since neighboring plots are known to be more alike in fertility than plots some distance removed, each replicate consists of a compact group of plots and is made approximately square in shape if feasible. Cultivations designed to keep the land clean of weeds will usually be carried out without regard to the replications, because it is not believed that results are affected by the order in which plots are cultivated. Similarly, the plots will generally be harvested in whatever order is most convenient. If, however, harvesting must be spread over a number of days, it is well to harvest the plots replication by replication, in case rainfall or other factors should produce changes in the weight of the crop from day to day. In other types of experimentation no real distinction might be made between replicates until relatively late in the experiment.

The principal advantages of randomized blocks are as follows.

1. By means of the grouping, more accurate results are usually obtained than with completely randomized designs.
2. Any number of treatments and any number of replicates may be included. With the design as described above, each treatment will have the same number of replicates. If extra replication is desired for some treatments, each of these may be applied to two units within every group. This device provides twice the standard number of replicates for the treatments in question, at the expense of some increase in the size of the group, which now contains more than a single replication. Similarly a treatment may be applied three or four times in a group.
3. The statistical analysis is straightforward. Mishaps which necessitate the omission of a complete group or of the entire data from one or more treatments do not introduce any complication in the analysis. When data from some individual units are lacking, the "missing-plot"

technique developed by Yates enables the available results to be fully utilized. Some extra computational labor is, however, involved, and if the gaps are numerous the design is less convenient in this respect than complete randomization.

4. If the experimental error variance is larger for some treatments than for others, an unbiased error for testing any specific combination of the treatment means can still be obtained.

No design is more frequently used than randomized blocks. Certainly if a satisfactory degree of precision is reached, there is little need to search for alternative designs.

It is worth noting that the replication means provide unbiased comparisons of the differences among replicates. Occasionally, these differences measure some property of the experimental material that is of interest. The variance-ratio test of the replications against the error m.s. requires some care in its interpretation. In a greenhouse experiment, for instance, a different type of soil might be used in each replication. Significant differences among replications might be due either to differences in soil type or to differences in the positions of the replications within the greenhouse.

4.22 Randomization. When the units have been grouped, the treatments are assigned at random to the units within each group. A new randomization is made for every group. Unless the number of treatments exceeds 16, tables 15.6 and 15.7 are convenient for this operation.

4.23 Statistical Analysis. The example comes from an experiment carried out by the North Carolina Agricultural Experiment Station at Rocky Mount, N. C., in 1944. The experiment tested the effects of 5 levels of application of potash, supplying respectively 36, 54, 72, 108, and 144 lb. K_2O per acre, on the yield and properties of cotton. The measure chosen for analysis is the Pressley strength index. This is found by measuring the breaking strength of a bundle of fibers of a given cross-sectional area. A single sample of cotton was taken from each plot, and 4 determinations were made on each sample. The figures in table 4.4 are the means of these 4 samples.* The experiment was arranged in 3 randomized blocks of 5 plots each.

To make the computing instructions more general we suppose that there are t treatments and r replicates, and that y denotes a typical observation.

* Since the machine which measures the index is calibrated in arbitrary units, no dimensions are ascribed to the data in table 4.4. The index can be converted approximately into pounds per square inch by means of a regression formula.

Step 1. Find the treatment totals (T_i), the replicate totals (R_j), and the grand total (G).

TABLE 4.4 STRENGTH INDEX OF COTTON IN A RANDOMIZED BLOCKS EXPERIMENT

Treatments Pounds K ₂ O per acre	Replications			Totals	Coded dressing
	1	2	3		
36	7.62	8.00	7.93	23.55	13
54	8.14	8.15	7.87	24.16	8
72	7.76	7.73	7.74	23.23	3
108	7.17	7.57	7.80	22.54	-7
144	7.46	7.68	7.21	22.35	-17
Totals	38.15	39.13	38.55	115.83	

Analysis of variance

Source of variation	d.f.	s.s.	m.s.	F
Replications	$(r - 1) = 2$	0.0971		
Treatments	$(t - 1) = 4$	0.7324	0.1831	4.19 *
Error	$(r - 1)(t - 1) = 8$	0.3495	0.0437	
Total	$rt - 1 = 14$	1.1790		

Subdivision of the treatments s.s.

	d.f.	s.s.	m.s.	F
Linear response	1	0.5663	0.5663	12.96**
Deviations	3	0.1661	0.0554	1.27

Step 2. The sums of squares are obtained as follows.

$$\text{Correction factor: } C = \frac{G^2}{tr} = \frac{(115.83)^2}{15} = 894.4393$$

$$\text{Total: } \sum y^2 - C = (7.62)^2 + (8.14)^2 + \dots + (7.21)^2 - 894.4393 = 1.1790$$

$$\text{Replications: } \sum \frac{R_j^2}{t} - C = \frac{(38.15)^2 + (39.13)^2 + (38.55)^2}{5} - 894.4393 = 0.0971$$

$$\text{Treatments: } \sum \frac{T_i^2}{r} - C = \frac{(23.55)^2 + (24.16)^2 + \dots + (22.35)^2}{3} - 894.4393 = 0.7324$$

$$\begin{aligned} \text{Error: (total s.s.)} - (\text{replications s.s.}) - (\text{treatments s.s.}) \\ = 1.1790 - 0.0971 - 0.7324 = 0.3495 \end{aligned}$$

The F -ratio for treatments (table 4.4) is 0.1831/0.0437, or 4.19, which is significant at the 5% level with 4 and 8 d.f. The treatment totals suggest that the strength decreases with increasing applications of potash, though there is a hint of a maximum in strength for the 54-lb.

application. Accordingly it is worth while to examine the shape of the response curve. We will first fit a linear regression on the amount of dressing. Note that successive increments of dressing are not equal, the last two increments being twice as large as the first two. This is a fairly common practice in cases where the effectiveness of an increase in dressing is expected to be smaller at the higher levels of application.

Although the regression can be calculated by the standard formula, a simple coding of the dressings lightens the work. If we place the lowest dressing (36 lb.) at zero and take 18 lb. as 1 unit on the scale, the dressings may be coded as 0, 1, 2, 4, and 6, respectively. A further device is to subtract a common amount from these values so that their mean is zero. Since the mean of the coded dressings is $1\frac{3}{5}$, this is the amount that must be subtracted. To avoid fractions, we multiply all values by 5 and subtract 13 from each product. The resulting coded dressings are shown in table 4.4; the signs have been changed so that the regression coefficient will be positive.

The regression coefficient is now obtained as

$$b = \frac{(13)(23.55) + (8)(24.16) + (3)(23.23) - (7)(22.54) - (17)(22.35)}{(3)[(13)^2 + (8)^2 + (3)^2 + (7)^2 + (17)^2]}$$

$$= \frac{31.39}{1740} = 0.0180$$

The factor (3) in the denominator is inserted to convert the treatment totals to means. The regression coefficient represents the average decrease in strength for a unit increase on the coded scale. Since 5 units on the coded scale correspond to 18 lb. K_2O , the decrease in strength for each additional 18 lb. K_2O is estimated as 0.090.

The contribution of the regression to the sum of squares in the analysis of variance is (by rule 1, section 3.42)

$$\frac{(31.39)^2}{1740} = 0.5663$$

as shown in the analysis of variance. The contribution from the linear regression is significant at the 1% level. Since the mean square for deviations from the regression, 0.0554, is only slightly above the error m.s., there seems no point in investigating a quadratic regression, though the reader may care to verify that its contribution is small.

The conclusion from the data is that increased dressings of potash produce a weaker fiber, the strength index declining by 0.090 for each 18 lb.-increment in K_2O . As an exercise we will obtain confidence limits for this rate of decline. We first require the estimated standard error for

the 0.090 figure. This can be found by expressing the figure as a linear function of the treatment means \bar{y}_i and applying rule 5a of section 3.51. We have

$$0.090 = \frac{5}{580}[13\bar{y}_1 + 8\bar{y}_2 + 3\bar{y}_3 - 7\bar{y}_4 - 17\bar{y}_5]$$

so that the estimated standard error is

$$\begin{aligned} \frac{s}{\sqrt{r}} \sqrt{\sum l_i^2} &= \frac{\sqrt{0.0437}}{\sqrt{3}} \frac{5}{580} \sqrt{(13)^2 + (8)^2 + \cdots + (17)^2} \\ &= 5 \frac{\sqrt{0.0437}}{\sqrt{1740}} = 0.0251 \end{aligned}$$

Alternatively, this value could be derived by noting that the square root of the F -value for the linear regression, i.e., $\sqrt{12.96}$, or 3.6, is the ratio of 0.090 to its estimated standard error. Thus the latter must be $0.09/3.6$, or 0.0250. Finally, for the 80% confidence limits, we have $(0.090 \pm 0.025 \times 1.397)$, or (0.090 ± 0.035) , where 1.397 is the value of t for a probability 0.20 and 8 d.f.

4.24 Standard Errors. The estimated standard error of the difference between two treatment means is

$$s_d = \sqrt{\frac{2s^2}{r}}$$

If some treatments receive extra replication, the general formula is

$$s_d = \sqrt{s^2 \left(\frac{1}{r_1} + \frac{1}{r_2} \right)}$$

In experiments where the error appears to be heterogeneous, a separate error may be obtained for any pair or for any group of treatments. For this a new randomized blocks analysis is carried out on those treatments which belong to the group under consideration.

4.25 Missing Data. The method of analysis when part of the data is missing is described by Yates (4.4). With a single missing unit, the first step is to calculate a value for the unit by means of the formula

$$y = \frac{rB + tT - G}{(r-1)(t-1)} \quad (4.1)$$

where B is the total of the remaining units in the block where the missing unit appears, T is the total of the yields of this treatment in the other blocks, and G is the grand total; r and t are the numbers of replicates and

treatments respectively. The analysis of variance is then carried out as usual except that 1 d.f. is subtracted from the total s.s. and from the error s.s.

The standard error of the difference between the mean of the treatment with a missing value and the mean of any other treatment is

$$\sqrt{s^2 \left[\frac{2}{r} + \frac{t}{r(r-1)(t-1)} \right]} \quad (4.2)$$

When there are several missing values, for units a, b, c, d, \dots , we first guess values by inspection for all units except a . Formula (4.1) is then used to find an approximation for a . With this approximation and the values previously assumed for c, d, \dots , we again use formula (4.1) to insert an approximation for b . After a complete cycle of these operations, a second approximation is found for a and so on until the new approximations are not materially different from those found previously. The analysis of variance is then completed; for each missing unit, 1 d.f. is subtracted from the total and error s.s.

Suppose that in the previous example two observations had been missing and that the data had appeared as follows.

Pounds K ₂ O	Replications			Totals
	1	2	3	
36	a	8.00	7.93	15.93
54	8.14	8.15	7.87	24.16
72	7.76	b	7.74	15.50
108	7.17	7.57	7.80	22.54
144	7.46	7.68	7.21	22.35
Totals	30.53	31.40	38.55	100.48

Since differences between replications are not pronounced, we might take as a trial value for a the mean of 8.00 and 7.93, or 7.96. For estimating b_1 , the first trial value of b , by formula (4.1) we now have

$$B = 31.40; \quad T = 15.50; \quad G = 100.48 + 7.96 = 108.44$$

so that

$$b_1 = \frac{(3)(31.40) + (5)(15.50) - 108.44}{8} = 7.91$$

For estimating a_2 , the second trial value of a , we now have

$$B = 30.53; \quad T = 15.93; \quad G = 100.48 + 7.91 = 108.39$$

so that

$$a_2 = \frac{(3)(30.53) + (5)(15.93) - 108.39}{8} = 7.86$$

Taking a_2 as 7.86, we find that the next trial value of b is 7.92. This is so close to the previous value that we may stop. Thus $a = 7.86$, $b = 7.92$.

The analysis of variance is now computed with these values inserted. There will be 12 d.f. in the total s.s., and 6 d.f. in the error s.s. The error m.s. is 0.0491.

To obtain a standard error for the comparison of two treatments, A and B , we assign an "effective" number of replicates to each treatment. Any replication of treatment A is counted as 1 when both A and B are present in the replicate, as $\frac{1}{2}$ when A is present but B is not, and as 0 when A is missing. The same rule is applied in scoring B . Suppose that in the example we are comparing the means of the 36- and 72-lb. dressings. The effective replication for each mean is scored as $1\frac{1}{2}$. The standard error of their difference is taken as $\sqrt{s^2(\frac{2}{3} + \frac{2}{3})}$, or $\sqrt{1.333s^2}$. If we are comparing the mean of the 36-lb. dressing with that of the 54-lb. dressing, which has no missing values, the score for the first mean is 2, and that for the second $2\frac{1}{2}$, so that the standard error of their difference is $\sqrt{s^2(\frac{1}{2} + \frac{2}{4})}$, or $\sqrt{0.9s^2}$.

This useful approximation is due to Yates; the exact formulae are laborious to compute. The approximation usually gives a standard error that is slightly too high. In the example the correct values for the standard errors of the two differences considered above are $1.069s$ and $0.937s$, as compared with the approximate values of $1.155s$ and $0.949s$, respectively.

As a result of the disturbance introduced by the missing units, the treatments m.s. is slightly too large; however, the variance-ratio test is unlikely to be much in error unless a substantial proportion of the units are missing. Yates (4.4) gives the method for obtaining an exact test.

4.26 Estimation of Efficiency. If E_b and E_e are the block and error m.s. and n_b , n_t , and n_e the block, treatment, and error degrees of freedom,

$$E_{e.r.} = \frac{n_b E_b + (n_t + n_e) E_e}{n_b + n_t + n_e} \quad (4.3)$$

is an estimate of the error variance of a completely randomized design with the same experimental material. Comparison of $E_{e.r.}$ and E_e , taking account of the change in numbers of error degrees of freedom (section 2.31), provides an estimate of the increase in accuracy which results from the grouping into replicates.

The result in (4.3) may be proved in various ways. Since there is no complete discussion of results of this type in the literature, one method of proof will be sketched; the details, which are a matter of algebraic manipulation, will not be given completely. This proof uses the proper-

ties of randomization. It is not the easiest proof, but requires very few assumptions.

Let the experiment have t treatments and r replications, and let e_i be the experimental error of the observation on the i th unit, and τ_j the effect of the j th treatment. No assumption is made about the nature or distribution of the errors. For this reason it is not necessary to introduce any specific symbol for the effect of the replication, since any type of effect can be represented by appropriate choice of the e 's. The treatment effect and the error are assumed to be additive; that is, if the randomization happens to put the j th treatment on the i th unit, their joint effect is $(\tau_j + e_i)$. Throughout the randomization the e_i are regarded as a set of fixed numbers, each associated with a specific unit.

Without loss of generality, we may assume that the total of all observations is zero, and that the totals of the e 's and the τ 's are both zero. Let

$$S = \sum_{i=1}^{rt} e_i^2; \quad T = \sum_{j=1}^t \tau_j^2$$

Further, for all possible randomizations of the randomized blocks design the replications will of course remain the same; consequently, for any given batch of data the replication totals remain unchanged. Let R denote the sum of squares of these totals, divided by t .

The main part of the proof consists in working out the average values of the mean squares in the analysis of variance, taken over all possible randomizations of each type of design. The results come out as shown in table 4.5.

TABLE 4.5 AVERAGE VALUES OF MEAN SQUARES TAKEN OVER THE RANDOMIZATION SETS

Completely randomized		
	d.f.	m.s.
Treatments	$(t - 1)$	$\frac{rT}{(t - 1)} + \frac{S}{(rt - 1)}$
Error	$t(r - 1)$	$\frac{S}{(rt - 1)}$
Randomized blocks		
	d.f.	m.s.
Replications	$(r - 1)$	$\frac{R}{(r - 1)}$
Treatments	$(t - 1)$	$\frac{rT}{(t - 1)} + \frac{S}{r(t - 1)} - \frac{R}{r(t - 1)}$
Error	$(r - 1)(t - 1)$	$\frac{S}{r(t - 1)} - \frac{R}{r(t - 1)}$

This analysis leads at once to the result. Since

$$n_b = (r - 1); \quad n_t + n_e = r(t - 1);$$

it is seen from table 4.5 that for randomized blocks the average value of $n_b E_b$ is R , while that of $(n_t + n_e) E_e$ is $(S - R)$. It follows that the average value of

$$\frac{n_b E_b + (n_t + n_e) E_e}{n_b + n_t + n_e} = \frac{R + S - R}{(rt - 1)} = \frac{S}{(rt - 1)}$$

But this is the average of the error m.s. for a completely randomized design on the same data.

As an illustration of the details, consider the treatments m.s. for randomized blocks, which is probably the hardest term. The square of the first treatment total is

$$T_1^2 = (r\tau_1 + e_1 + e_2 + \cdots + e_r)^2$$

When we average over all possible randomizations, there is no contribution from terms in τe_i , since the treatment appears equally often on all units, and the total of the e 's is zero. The average value of the contribution from the r terms in e^2 is the sum of the squares of *all* the e 's, divided by t , or S/t , since the treatment appears on any specific unit in a fraction $1/t$ of all randomizations.

The contribution from terms in $e_i e_j$ may be written

$$e_1(e_2 + e_3 + \cdots + e_r) + e_2(e_1 + e_3 + \cdots + e_r) + \cdots + e_r(e_1 + e_2 + \cdots + e_{r-1}) \quad (4.4)$$

The randomization is restricted so that every e comes from a different replication. If the subscript denotes the replicate, the mean value of $e_i e_j$ is $R_i R_j / t^2$, where R_i is the replication total of the e 's. Consequently, the mean value of (4.4) is

$$\begin{aligned} & \frac{1}{t^2} [R_1(R_2 + R_3 + \cdots + R_r) + R_2(R_1 + R_3 + \cdots + R_r) + \cdots \\ & + R_r(R_1 + R_2 + \cdots + R_{r-1})] = -\frac{1}{t^2} (R_1^2 + R_2^2 + \cdots + R_r^2) = -\frac{R}{t} \end{aligned}$$

since the total of the R_i is zero, and since R will be recognized as the sum of squares of the R_i , divided by t . Hence

$$E(T_1^2) = r^2 \tau_1^2 + \frac{S}{t} - \frac{R}{t}$$

If this expression is summed for all t treatments and divided by $r(t - 1)$ to give the treatments m.s., the result in table 4.5 follows.

4.3 Double Grouping: Latin Squares

4.31 Description. In the latin square the treatments are grouped into replicates in two different ways. Examples of latin squares are shown for different numbers of treatments in plan 4.1 (p. 119). It will be noted that every row and every column of any square is a complete replication. The effect of the double grouping is to eliminate from the errors all differences among rows and equally all differences among columns. Thus the latin square provides more opportunity than randomized blocks for the reduction of errors by skillful planning.

The experimental material should be arranged and the experiment conducted so that the differences among rows and columns represent major sources of variation. Some examples of the uses of latin squares in various fields of research may indicate the utility of the design. In field experiments the plots are usually laid out in a square formation, so that soil fertility and other variations in two directions are controlled. Variations along and across the greenhouse bench may be similarly handled in greenhouse experiments. Occasionally the latin square is advantageous even when the plots form a continuous line. In this case the rows may be compact blocks of land while the columns specify the order within each block. If the yield gradient is suspected to be in the same direction all along the line, blocks and order in blocks together remove the effects of the gradient more thoroughly than a single control. An example of this type is shown below.

TABLE 4.6 A LATIN SQUARE WITH THE PLOTS IN ONE CONTINUOUS LINE

Seven treatments A, B, C, D, E, F, G

(Experimental limitations force the plot units to go at right angles to gradient.)

Rep. I	II	III	IV	V	VI	VII
A	D	G	B	E	C	F
G	F	B	E	C	A	D
C	F	B	E	C	A	D
F	B	E	C	A	D	G
B	C	A	D	G	F	E
E	D	G	F	E	C	A
D	G	F	E	C	A	B
C	A	B	D	G	F	E
A	B	D	G	F	E	C
B	D	G	F	E	C	A
D	G	F	E	C	A	B
G	F	E	C	A	B	D
F	E	C	A	B	D	G
E	C	A	B	D	G	F
C	A	B	D	G	F	E
A	B	D	G	F	E	C

Main and Tippet (4.5) describe how 4×4 latin squares were employed in experiments on the weaving of cotton cloth. The purpose of a series of experiments was to investigate the effect of the sizing treatment applied to the warp. The criterion was the number of breaks in the warp during weaving. Four warps, each with a different sizing treatment, were woven simultaneously on 4 different looms, which could be supervised by a single weaver. Then each warp was moved to a different loom of the set so that after 4 periods every sizing treatment had been tested on all 4 looms.

If *A*, *B*, *C*, *D* represent the 4 warps, the latin square used was as follows.

Periods	Looms			
	1	2	3	4
I	<i>A</i>	<i>D</i>	<i>B</i>	<i>C</i>
II	<i>D</i>	<i>C</i>	<i>A</i>	<i>B</i>
III	<i>C</i>	<i>B</i>	<i>D</i>	<i>A</i>
IV	<i>B</i>	<i>A</i>	<i>C</i>	<i>D</i>

This arrangement eliminates constant differences among the looms, which were found to be large, and also differences among the 4 periods of weaving. No "period" differences were apparent—a result which might be anticipated, the authors suggest, because humidity and warp tension were controlled throughout the experiment.

Frequently, particularly in industry, an experiment requires a series of operations each of which may introduce variability into the final results. In such cases the latin square may be useful in a preliminary investigation of the sources of variation. For example, in the preparation of an explosive mixture used in primers, variation may occur either in the mixing of the ingredients of the explosive or in the process of charging. One experiment of this type involved 4 mixing-blending teams and 4 charging operators. On each day, the product of each team was sent to a different charging operator, the arrangement being changed daily according to the following 4×4 latin square (letters *W*, *X*, *Y*, and *Z* represent the mixing-blending teams).

	Charging operators			
	1	2	3	4
Monday	<i>W</i>	<i>Z</i>	<i>X</i>	<i>Y</i>
Tuesday	<i>X</i>	<i>W</i>	<i>Y</i>	<i>Z</i>
Wednesday	<i>Y</i>	<i>X</i>	<i>Z</i>	<i>W</i>
Thursday	<i>Z</i>	<i>Y</i>	<i>W</i>	<i>X</i>

The latin square analysis of variance enables us to isolate consistent differences amongst the teams and consistent differences amongst the chargers, as well as day-to-day variations.

To quote a more complex example, a 12×12 latin square was used by Chen, Bliss, and Robbins (4.6) to estimate the toxicities of 12 poisons when applied to cats. A single replication of this experiment occupied a complete day, and the experiment was repeated for 12 days. Each drug was injected into the femoral vein of a cat at 1 cc. per minute until the cat died, and the measurement recorded was the dose required to cause death. An observer could administer only two of the drugs at a

given time. Consequently, three observers were used, each treating two cats in the morning and two in the afternoon. The rows of the latin square were used to eliminate systematic differences between observers and between the morning and afternoon injections. The columns represent days. The experimental plan was as follows (letters represent drugs).

Time	Observer	Day											
		1	2	3	4	5	6	7	8	9	10	11	12
10:30 A.M.	I	<i>I</i>	<i>J</i>	<i>B</i>	<i>L</i>	<i>H</i>	<i>G</i>	<i>F</i>	<i>K</i>	<i>D</i>	<i>E</i>	<i>A</i>	<i>C</i>
		<i>K</i>	<i>G</i>	<i>J</i>	<i>H</i>	<i>I</i>	<i>B</i>	<i>L</i>	<i>C</i>	<i>E</i>	<i>F</i>	<i>D</i>	<i>A</i>
	II	<i>B</i>	<i>L</i>	<i>G</i>	<i>C</i>	<i>D</i>	<i>J</i>	<i>K</i>	<i>E</i>	<i>H</i>	<i>A</i>	<i>F</i>	<i>I</i>
		<i>E</i>	<i>D</i>	<i>F</i>	<i>G</i>	<i>J</i>	<i>K</i>	<i>A</i>	<i>L</i>	<i>C</i>	<i>I</i>	<i>B</i>	<i>H</i>
	III	<i>C</i>	<i>K</i>	<i>A</i>	<i>B</i>	<i>F</i>	<i>L</i>	<i>I</i>	<i>D</i>	<i>G</i>	<i>H</i>	<i>J</i>	<i>E</i>
		<i>F</i>	<i>H</i>	<i>K</i>	<i>E</i>	<i>G</i>	<i>C</i>	<i>D</i>	<i>B</i>	<i>A</i>	<i>L</i>	<i>I</i>	<i>J</i>
2:30 P.M.	I	<i>J</i>	<i>C</i>	<i>E</i>	<i>K</i>	<i>A</i>	<i>I</i>	<i>H</i>	<i>F</i>	<i>B</i>	<i>G</i>	<i>L</i>	<i>D</i>
		<i>D</i>	<i>F</i>	<i>I</i>	<i>A</i>	<i>L</i>	<i>E</i>	<i>C</i>	<i>G</i>	<i>J</i>	<i>B</i>	<i>H</i>	<i>K</i>
	II	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	<i>G</i>	<i>H</i>	<i>I</i>	<i>J</i>	<i>K</i>	<i>L</i>
		<i>H</i>	<i>E</i>	<i>L</i>	<i>J</i>	<i>C</i>	<i>A</i>	<i>B</i>	<i>I</i>	<i>K</i>	<i>D</i>	<i>G</i>	<i>F</i>
	III	<i>G</i>	<i>I</i>	<i>D</i>	<i>F</i>	<i>K</i>	<i>H</i>	<i>J</i>	<i>A</i>	<i>L</i>	<i>C</i>	<i>E</i>	<i>B</i>
		<i>L</i>	<i>A</i>	<i>H</i>	<i>I</i>	<i>B</i>	<i>D</i>	<i>E</i>	<i>J</i>	<i>F</i>	<i>K</i>	<i>C</i>	<i>G</i>

During the course of the experiment every drug was given equally often by each observer and equally often in the morning and afternoon. The arrangement is a good example of the way in which a single grouping (rows) can control simultaneously more than one potential source of variation. All three major variables—day, time of day, and observer—would have inflated the experimental error if left uncontrolled.

4.32 Number of Replications. The chief restriction on the utility of the latin square is that the number of replicates equals the number of treatments; if the latter is considerable, the number of replications required becomes impractical. Squares larger than 12×12 are seldom used, while the most common range is from the 5×5 to the 8×8 square. Latin squares also suffer to some extent from the same dis-

advantage as randomized blocks in that the experimental error per unit is likely to increase with the size of square.

The small squares provide only a few degrees of freedom for the estimation of error—none with the 2×2 , two with the 3×3 , and six with the 4×4 . This fact precludes the use of single 2×2 squares, while the 3×3 and 4×4 squares must produce a substantial reduction in error over randomized blocks or complete randomization to counterbalance the loss of degrees of freedom. More than one square may, however, be included in the same experiment. Three 3×3 squares (9 replicates) furnish 10 error d.f., while two 4×4 squares give 15 d.f., provided that in each case the "squares \times treatments" interaction can be pooled with error. A considerable number of 2×2 squares would be required, since the error degrees of freedom are 1 less than the number of squares.

4.33 Randomization. A complete representation of the squares from 4×4 to 6×6 and sample squares up to the 12×12 is given by Fisher and Yates (4.3). For squares up to the 6×6 , the randomization procedure given in this reference selects (with a minimum of labor) a square at random from all latin squares of a given size. For a discussion of the theoretical basis of the randomization, see Yates (4.7).

Examples of latin squares are shown in plan 4.1 (p. 119). The method of randomization for plan 4.1 is as follows.

3×3 . Arrange the columns at random and the last 2 rows at random.

4×4 . Select at random one of the 4 squares. Arrange at random all columns and the last 3 rows. It is equally good, though not strictly necessary, to randomize *all* rows and columns.

5×5 and higher squares. Arrange all rows, columns, and treatments independently at random.

For 3×3 and 4×4 squares, this procedure selects one square at random from all possible squares. For 5×5 and larger squares, some types of squares have no chance of being selected if plan 4.1 is used. Unless latin squares are used very frequently, however, the randomization sets are sufficiently large for experimental plans.

4.34 Statistical Analysis. Several experiments have demonstrated that people find it difficult to select, by personal judgment, unbiased samples even from relatively small populations that can be thoroughly inspected before selection. In this experiment, each population consisted of a small area of wheat, containing about 80 shoots, the shoots being slightly over 2 feet high. There were 12 samplers, all experienced in studying the growth of wheat. Each sampler inspected each area and measured the heights of 8 shoots whose heights were to give a represent-

ative sample of the shoot heights in the area. The quantity that will be analyzed is the difference between the mean height of the 8 selected shoots and the true mean height in the corresponding area, in other words the sampler's error.

The samplers were divided into two groups of six, of which we will consider only the first group. The samplers represent the experimental treatments. There were 6 areas, each sampled by all 6 samplers in the group. These areas, which serve as the replications, may be taken as the columns of the 6×6 latin square. The rows of the square prescribed the order in which each man sampled the 6 areas (see table 4.7). This

TABLE 4.7 SAMPLER'S ERRORS IN SHOOT HEIGHTS (CENTIMETERS)
 6×6 LATIN SQUARE

Order	Areas						Totals
	1	2	3	4	5	6	
I	F+ 3.5	B+ 4.2	A+6.7	D+6.6	C+4.1	E+3.8	+28.9
II	B+ 8.9	F+ 1.9	D+5.8	A+4.5	E+2.4	C+5.8	+29.3
III	C+ 9.6	E+ 3.7	F-2.7	B+3.7	D+6.0	A+7.0	+27.3
IV	D+10.5	C+10.2	B+4.6	E+3.7	A+5.1	F+3.8	+37.9
V	E+ 3.1	A+ 7.2	C+4.0	F-3.3	B+3.5	D+5.0	+19.5
VI	A+ 5.9	D+ 7.6	E-0.7	C+3.0	F+4.0	B+8.6	+28.4
Totals	+41.5	+34.8	+17.7	+18.2	+25.1	+34.0	+171.3 = G
Totals for samplers							
	<u>A</u>	<u>B</u>	<u>C</u>	<u>D</u>	<u>E</u>	<u>F</u>	
	+36.4	+33.5	+36.7	+41.5	+16.0	+7.2	

feature served two purposes. It helped to prevent the men from getting in each other's way, since no two men worked on the same area at the same time, and it permitted an examination of the effects of the order of sampling on the errors.

Since only 3 of the 36 errors in table 4.7 are negative, there is evidence of a rather consistent tendency towards overestimation of the shoot heights. For simplicity, the observations will be referred to as measures of bias, though of course they contain a component due to chance fluctuations. There is also an indication, on casual inspection of the table, that the bias varies from one sampler to another. This point will be tested in the analysis of variance. The computations are as follows. First find the totals for each row, column, and latin letter and the grand

total, G , shown in table 4.7. The following sums of squares are then calculated.

$$\text{Correction factor: } C = \frac{G^2}{r^2} = \frac{(171.3)^2}{36} = 815.10$$

$$\text{Total: } (3.5)^2 + (4.2)^2 + \cdots + (8.6)^2 - C = 1144.73 - 815.10 = 329.63$$

$$\text{Rows: } \frac{1}{6}[(28.9)^2 + \cdots + (28.4)^2] - C = 843.70 - 815.10 = 28.60$$

$$\text{Columns: } \frac{1}{6}[(41.5)^2 + \cdots + (34.0)^2] - C = 893.97 - 815.10 = 78.87$$

$$\text{Treatments: } \frac{1}{6}[(36.4)^2 + \cdots + (7.2)^2] - C = 970.70 - 815.10 = 155.60$$

The divisor 6 is replaced by r for an $r \times r$ square.

These sums of squares are entered in the analysis of variance. The error s.s. is found by subtraction.

TABLE 4.8 ANALYSIS OF VARIANCE FOR A 6×6 LATIN SQUARE

Source of variation	d.f.	s.s.	m.s.	F
Rows (order of sampling)	$(r - 1) = 5$	28.60	5.720	1.72
Columns (areas)	$(r - 1) = 5$	78.87	15.774	4.74 **
Treatments (samplers)	$(r - 1) = 5$	155.60	31.120	9.35 **
Error	$(r - 1)(r - 2) = 20$	66.56	3.328	
Total	$(r^2 - 1) = 35$	329.63		

To obtain the mean squares, each sum of squares is divided by the corresponding number of degrees of freedom. The F value for samplers is $31.120/3.328$, or 9.35, well beyond the 1% level, which is 4.10 for 5 and 20 d.f. This shows that the extent of the overestimation varies from one sampler to another. The means for the samplers are given below.

A	B	C	D	E	F
+6.07	+5.58	+6.12	+6.92	+2.67	+1.20

The estimated standard error of each mean is $\sqrt{s^2/r}$, where s^2 is the error m.s., or in this case $\sqrt{(3.328)/6} = 0.745$. For testing the difference between a pair of means, the standard error is $\sqrt{2} (0.745) = 1.053$. Since the 5% t value for 20 d.f. is 2.086, the difference between two means must be at least $(2.086)(1.053)$, or 2.20, in order to attain significance at this level. It appears that the samplers fall into two sets— A, B, C, D and E, F , the biases being significantly smaller for E and F than for the others.

Some further information about the nature of the biases can be obtained from an examination of the row and column totals. From table

4.8 the F value for rows is seen to be 1.72, which is below the 5% value, 2.71. No clear effect of the order in which the areas were sampled has therefore been established; further, the row totals (table 4.7) do not suggest any consistent trend.

On the other hand, the F value for the columns m.s., 4.74, is significant at the 1% level, so that the bias varies from area to area. Previous observations had indicated that the bias might depend on the mean shoot height of the area. The column totals are ranked below in order of increasing true mean shoot height.

Area	True mean shoot height (centimeters)	Column total (centimeters)
1	59.0	+41.5
2	66.2	+34.8
6	72.3	+34.0
4	74.5	+18.2
5	76.0	+25.1
3	76.4	+17.7

Apparently on areas where the shoots are not so high the positive bias is greater than on areas with relatively high shoots. The result is not surprising, since the sampler had to stoop in order to do his work. The linear regression of the column totals on the true mean heights contributes 59.88 to the sum of squares for columns, 78.87. The mean square for deviations from this regression (4 d.f.) is only 4.75, which is not significantly greater than the error m.s. The inference is that the large value of the mean square for columns is due mainly to the negative correlation between the bias and the mean height. For further discussion of this experiment, see (4.8).

The analysis of two 4×4 squares is discussed in (4.9). A numerical example with five 3×3 squares will be found in (4.10). It should be noted that with several latin squares the interaction of treatments with squares may be tested. If there is no reason to expect that this interaction is real, the corresponding sum of squares may be combined with error.

4.35 Standard Errors. The formula $\sqrt{\frac{2s^2}{r}}$ gives the estimated standard error of the difference between two treatment means, where s^2 is the mean square per unit and r the number of replicates. In cases where heterogeneity of errors is suspected, the error cannot be subdivided with the same ease as in randomized blocks. It is possible to remove from the error s.s. the contribution of any one of the *treatments*, Yates (4.11). For

each yield y_i of the treatment, calculate the quantity

$$d_i = R_i + C_i - ry_i$$

where R_i and C_i are the corresponding row and column totals, and r is the number of treatments. The sum of squares of deviations of the quantities d_i , divided by $r(r-2)$, is the contribution of this treatment to the error s.s. and has $(r-1)$ degrees of freedom. For illustration, suppose that we wish to calculate the contribution of sampler F to the error in section 4.34. From table 4.7 the first d value is

$$d_1 = (+28.9) + (+41.5) - 6(+3.5) = +49.4$$

and the remaining values will be found to be +52.7, +61.2, +57.5, +29.5, +49.1, with a total +299.4. The sum of squares of deviations from the mean is 610.34. Divided by 24, this gives 25.43, with 5 d.f., for the contribution of sampler F to the error s.s. Consequently the error s.s. (table 4.8) may be divided as shown below.

	d.f.	s.s.	m.s.
Total error	20	66.56	3.328
Contribution of F	5	25.43	5.086
Remainder	15	41.13	2.742

The error m.s. for the suspected treatment may be compared with the mean square for the remaining $(r-1)(r-3)$ degrees of freedom, to test whether the treatment shows abnormally high variation. In this example the variance ratio is 1.85, with 5 and 15 d.f. If it is concluded that the treatment is unusually variable, the mean square with $(r-1)(r-3)$ degrees of freedom is used as error for the other treatments. It should be stressed that the ordinary variance-ratio tables can be used in this test only if the decision to make the test was taken before examining the individual results. If the test is made merely because one treatment looks anomalous on inspection, different significance levels that take account of this fact are needed, as discussed in section 3.53.

4.36 Missing Values. Except for changes in the formulae, the procedure is similar to that for randomized blocks (section 4.25). A single missing value is substituted by means of the formula, reference (4.4)

$$y = \frac{r(R + C + T) - 2G}{(r-1)(r-2)} \quad (4.5)$$

where R , C , T , and G are respectively the totals of the row, column, and treatment which contain the missing value and the grand total. The

estimated standard error of the difference between the corresponding treatment mean and the mean of a treatment with no missing values is

$$\sqrt{s^2 \left(\frac{2}{r} + \frac{1}{(r-1)(r-2)} \right)} \quad (4.6)$$

When several values are missing, repeated application is made of formula (4.5) as described in reference (4.4). When values have been inserted for all missing data, the usual analysis of variance is calculated. One degree of freedom is subtracted from the error d.f. for each missing value.

The exact formula for the standard error of a treatment mean is complex. Yates (4.4) proposed a useful approximate rule, which shows the number of replicates to be assigned to any treatment mean for a comparison with another treatment mean. Each observation on one treatment is given one replication if the other treatment is present in both the corresponding row and column. The replication is $\frac{2}{3}$ when the other treatment is missing in the row *or* column and is $\frac{1}{3}$ when the other treatment is missing in both the row and column. When the treatment itself is missing, the replication is 0. Although complicated at first sight, the rule is not difficult to apply and is reasonably accurate. Consider for instance the following 6×6 square with 3 units missing (two *B*'s and one *E*) as indicated by the parentheses ().

<i>B</i>	<i>E</i>	<i>C</i>	<i>F</i>	<i>D</i>	<i>A</i>
<i>F</i>	<i>D</i>	(<i>B</i>)	<i>E</i>	<i>A</i>	<i>C</i>
<i>C</i>	<i>A</i>	<i>E</i>	<i>D</i>	<i>B</i>	<i>F</i>
(<i>E</i>)	<i>C</i>	<i>A</i>	(<i>B</i>)	<i>F</i>	<i>D</i>
<i>A</i>	<i>F</i>	<i>D</i>	<i>C</i>	<i>E</i>	<i>B</i>
<i>D</i>	<i>B</i>	<i>F</i>	<i>A</i>	<i>C</i>	<i>E</i>

Taking the yields of *B* and *E* in the order of the rows, we ascribe the following numbers of replicates.

$$B: \frac{2}{3} + 0 + 1 + 0 + 1 + 1 = 3\frac{2}{3} = \frac{11}{3}$$

$$E: 1 + \frac{1}{3} + \frac{2}{3} + 0 + 1 + 1 = 4$$

Hence the standard error of the difference between the treatment means is $\sqrt{s^2(\frac{3}{11} + \frac{1}{4})}$. For the difference between *A* and *B*, the reader may verify that the standard error is $\sqrt{s^2(\frac{3}{14} + \frac{1}{4})}$. Notice that the effective replication for a treatment may change from one comparison to another.

The analysis required when a single row, column, or treatment is missing is described by Yates (4.11). When more than one row, column, or

treatment is missing, consult Yates and Hale (4.12). These methods are also presented by DeLury (4.13).

4.37 Estimation of Efficiency. The effectiveness of either the row or the column grouping may be tested from the results of latin squares. The expression

$$E' = \frac{n_r E_r + (n_t + n_e) E_e}{n_r + n_t + n_e}$$

is an estimate of the error m.s. which would have been obtained if the row grouping had not been used, i.e., if the design had been randomized blocks with the columns as blocks. In the formula, E_r and E_e are the mean squares for rows and error, respectively, and n_r , n_t , n_e are the numbers of degrees of freedom for rows, treatments, and error. The increased number of degrees of freedom with randomized blocks is taken into account by the method of section 2.31. The effectiveness of the column grouping is tested similarly.

A 5×5 experiment on potatoes will serve as an illustration. The mean squares for the yields of potatoes were as follows.

	d.f.	m.s.
Rows	4	62.6
Columns	4	104.7
Treatments	4	272.0
Error	12	26.2

If the experiment had been conducted in randomized blocks with the columns as blocks, the estimated error m.s. would be

$$\frac{4(62.6) + 16(26.2)}{20} = 33.5$$

The degrees of freedom available for error would increase from 12 to 16. To take account of the advantage of the additional degrees of freedom (section 2.31), we reduce 33.5 by the multiplier $(13)(19)/(17)(15)$, or 0.969. This gives 32.5 as a comparable mean square for randomized blocks. The row grouping increased the information by an estimated 25%. Of course a considerable number of these comparisons would be needed for drawing general conclusions.

4.4 Cross-over Designs

4.41 Description. In dairy husbandry and biological assay a design has been used which closely resembles the latin square but may have some advantages when the number of treatments is small. In the sim-

plest case where there are two treatments *A* and *B*, the units are first grouped into pairs as if a randomized block design were to be used. Suppose that from previous knowledge one unit in each pair is expected to give a higher response than the other and that the difference in favor of the superior unit is expected to be about the same in all pairs. It will clearly be advisable to ensure that each treatment is applied to the "better" member in half the replicates and to the "poorer" member in the other half. The pairs or replicates, which must be even in number, are divided *at random* into two equal sets, the first set to receive treatment *A* on the superior member of each pair, the second to receive *B*.

The experiment discussed in chapter 1 provides an example. The data (shown in table 4.10) are the times required to compute the sums of squares of 27 observations on each of two machines whose speeds it was desired to compare. Ten sums of squares were calculated, making 10 replications of the experiment. The cross-over design was used because it was thought that the second computation of a sum of squares might be faster than the first, so that for a fair comparison each machine should be used first in five of the ten replications. The randomization assigned machine *A* to be used first in replications 1, 3, 6, 8, and 9.

The degrees of freedom in the analysis of variance are as follows.

	d.f.
Columns (pairs or replicates)	9
Rows	1
Treatments	1
Error	8
	<hr/>
Total	19

Note that only 1 d.f. is assigned to rows. That is, the arrangement removes from error only the *average* difference between the 2 rows. In so far as the difference is not constant but varies from one replication to another, this variation enters into the experimental error.

In some experiments it is known that this difference *will* vary, and it is possible to estimate in advance (at least roughly) whether the difference will be large or small for a given replicate. Suppose that the difference is expected to decrease steadily from replicate 1 to replicate 10. With this knowledge a more accurate design is obtained by the use of five 2×2 latin squares, as shown below.

TABLE 4.9 FIVE 2×2 LATIN SQUARES FOR 2 TREATMENTS IN 10 REPLICATES

	Square									
Rows	I		II		III		IV		V	
	1	2	3	4	5	6	7	8	9	10
Better	<i>A</i>	<i>B</i>	<i>A</i>	<i>B</i>	<i>B</i>	<i>A</i>	<i>A</i>	<i>B</i>	<i>A</i>	<i>B</i>
Poorer	<i>B</i>	<i>A</i>	<i>B</i>	<i>A</i>	<i>A</i>	<i>B</i>	<i>B</i>	<i>A</i>	<i>B</i>	<i>A</i>

In this arrangement we remove from error 5 d.f. for rows, one in each square. Thus not only the average difference is removed, but also the variation in the difference from square to square. The combined analysis of variance of the 5 squares is set out as follows.

	d.f.
Squares	4
Columns within squares	5
Rows within squares	5
Treatments	1
Error	4
	—
Total	19

The 9 d.f. for squares and columns within squares are exactly the same as the 9 d.f. for columns in the cross-over design. Owing to the more complete elimination of the row effects, there are only 4 d.f. for error instead of 8 with the cross-over plan.

To summarize, the cross-over design is particularly appropriate when the difference between the rows is substantially the same in all replicates, for in this case the whole of the real difference between rows is concentrated in the single degree of freedom and the error variance is no larger than that of the latin squares. Even if the difference between rows is known to be variable, the cross-over design may be preferable in small experiments where few degrees of freedom are available for error.

In dairy husbandry the cross-over may be used to compare the effects of two feeding rations on the amount and quality of milk produced by the cow. Since cows vary greatly in their milk production, each ration is tested on every cow by feeding it during either the first or the second half of the period of lactation, so that each cow gives a separate replicate. The milk yield of a cow declines sharply from the first to the second half of its period, so that the first half is always "better," in the sense above. Whether a cross-over is superior to a set of latin squares depends on circumstances. The rate of decline is not constant from cow to cow; it is greater in general for high-yielding than for low-yielding cows. Thus if previous production records are available, the cows may be divided into pairs on the basis of yielding ability, each pair being made a separate 2×2 latin square. This plan is likely to give a smaller error than the cross-over, though sometimes not sufficiently smaller to counterbalance the loss of degrees of freedom.

The design can be used with any number of treatments, subject to the restriction that the number of replicates must be a multiple of the number of treatments. With three treatments, for example, a plan can be

drawn up from the 3 cycles ABC , BCA , CAB , where the order of the letters denotes the row to which each treatment is applied. Each cycle is allotted at random to one-third of the replicates. For higher numbers of treatments a design is constructed in the same way from the columns of any latin square. When the number of treatments exceeds four, however, the degrees of freedom for error are sufficiently large so that a set of latin squares is usually preferable.

4.42 Statistical Analysis. All sums of squares are calculated in the usual way. For the example (table 4.10), the computations are as follows.

$$\begin{aligned}
 \text{Correction factor: } & \frac{(352)^2}{20} & = 6195.2 \\
 \text{Total: } & (30)^2 + (14)^2 + (21)^2 + \dots + (23)^2 - 6195.2 & = 910.8 \\
 \text{Columns: } & \frac{1}{2}[(44)^2 + (42)^2 + \dots + (47)^2] - 6195.2 & = 357.8 \\
 \text{Rows: } & \frac{(194 - 158)^2}{20} & = 64.8 \\
 \text{Treatments: } & \frac{(216 - 136)^2}{20} & = 320.0
 \end{aligned}$$

TABLE 4.10 CROSS-OVER EXPERIMENT FOR COMPARING THE SPEEDS OF TWO COMPUTING MACHINES *A* AND *B*

Time (seconds minus 2 minutes) taken to calculate a sum of squares

Rows	Columns (replications)										Totals
	1	2	3	4	5	6	7	8	9	10	
First (poorer)	A30	B21	A22	B13	B13	A29	B 7	A12	A23	B24	194
Second (better)	B14	A21	B 5	A22	A18	B17	A16	B14	B 8	A23	158
Totals	44	42	27	35	31	46	23	26	31	47	352

Treatment totals: *A* 216; *B* 136

Analysis of variance

Source of variation	d.f.	s.s.	m.s.
Columns (replications)	9	357.8	39.8
Rows (first versus second)	1	64.8	64.8
Treatments	1	320.0	320.0 **
Error	8	168.2	21.0
Total	19	910.8	

The error s.s. is found by subtraction. The F -ratio for treatments is significant at the 1% level.

We will calculate the confidence limits for the true difference in speed which were quoted in section 1.2. The average observed difference is 8.0 seconds in favor of machine B . The estimated standard error of this difference is $\sqrt{2(21.0)/10}$, or 2.049, based on 8 d.f. Consequently the confidence limits for the true difference are

$$8.0 \pm 2.049t'$$

where t' is the value of Student's t corresponding to $(1 - P)$, where P is the confidence probability chosen. For $P = .95$, $t' = 2.306$, and the limits are (8.0 ± 4.7) , or 3.3 and 12.7. For $P = .80$, $t' = 1.397$, and, for $P = .99$, $t' = 3.355$, giving confidence limits of (5.1, 10.9) and (1.1, 14.9), respectively, as quoted.

Worked examples for the case where there are two treatments are also given by Brandt (4.14) and Fieller (4.15); the latter also illustrates the analysis of covariance for this design and the procedure when one column is missing.

In general, if there are t treatments and r replicates, the degrees of freedom subdivide as follows.

	d.f.
Columns	$(r - 1)$
Rows	$(t - 1)$
Treatments	$(t - 1)$
Error	$(t - 1)(r - 2)$
Total	$(tr - 1)$

4.43 Standard Errors. The usual formula $\sqrt{\frac{2s^2}{r}}$ holds for the standard error of the difference between two treatment means.

The substitution formula for a missing value is

$$y = \frac{rC + r(R + T) - 2G}{(t - 1)(r - 2)} \quad (4.7)$$

where the capital letters refer respectively to the totals of the column, row, and treatment in which the missing value appears and to the grand total. The difference between the mean of the affected treatment and the mean of a treatment with no missing value has a standard error

$$\sqrt{s^2 \left[\frac{2}{r} + \frac{t}{r(t - 1)(r - 2)} \right]} \quad (4.8)$$

4.5 Triple Grouping: Graeco-latin Squares

4.51 Description. In this arrangement the treatments are grouped into replicates in three different ways with the consequence that the effects of three different sources of variation are equalized for all treatments.

TABLE 4.11 A 5×5 GRAECO-LATIN SQUARE

Litters	Pen (weight groups)				
	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>
I	A_1	B_3	C_5	D_2	E_4
II	B_2	C_4	D_1	E_3	A_5
III	C_3	D_5	E_2	A_4	B_1
IV	D_4	E_1	A_3	B_5	C_2
V	E_5	A_2	B_4	C_1	D_3

The additional grouping (usually represented by greek letters) is denoted here by subscripts. Each treatment (A, \dots, E) appears once in each row and column and once with each subscript ($1 \dots 5$).

One example is an arrangement proposed by Dunlop (4.16) for testing 5 feeding treatments on pigs. The arrangement requires 5 pigs from each of 5 litters (I, II, III, IV, and V), the effects of litter differences being equalized by making litters correspond to the rows of the square. The animals are housed in 5 pens (a, b, c, d , and e) which constitute the columns. The columns are used also to control differences in weight; i.e., the heaviest animal in each litter is assigned to the first pen, the next heaviest to the second pen, and so on; consequently the columns eliminate simultaneously the pen differences and the principal variations in weight within litters. Thus far the design is an ordinary latin square.

The subscripts signify the positions of the five feeding crates within each pen. This extra control, though possibly unnecessary, was suggested for several reasons. The first and fifth crates in each pen were of different construction from the other crates; moreover the pigs occupying the end crates have less "company" when feeding.

The graeco-latin square has not been used often, probably because the units can seldom be balanced conveniently in all three groupings. In the example discussed above, the practical difficulty was to secure five litters of which each contained five or more pigs. An interesting industrial application is described by Tippet (4.17).

The designs have been constructed for all numbers of treatments from 3 to 12, except 6 and 10; examples are shown in plan 4.2 (p. 120).

4.52 Randomization. Arrange the rows and columns independently at random. Assign the latin letters and the subscripts at random to their respective classifications.

4.53 Statistical Analysis. The sums of squares for rows, columns, treatments, and subscripts are all obtained in the usual way. For an $r \times r$ square the error has $(r - 1)(r - 3)$ degrees of freedom. This number is rather inadequate when r is less than 6.

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PLANS

Plan 4.1

Selected latin squares

 3×3

A	B	C
B	C	A
C	A	B

1	A	B	C	D
B	A	D	C	
C	D	B	A	
D	C	A	B	

 4×4

2	A	B	C	D
B	C	D	A	
C	D	A	B	
D	A	B	C	

3	A	B	C	D
B	D	A	C	
C	A	D	B	
D	C	B	A	

4	A	B	C	D
B	A	D	C	
C	D	A	B	
D	C	B	A	

 5×5

A	B	C	D	E
B	A	E	C	D
C	D	A	E	B
D	E	B	A	C
E	C	D	B	A

 6×6

A	B	C	D	E	F
B	F	D	C	A	E
C	D	E	F	B	A
D	A	F	E	C	B
E	C	A	B	F	D
F	E	B	A	D	C

 7×7

A	B	C	D	E	F	G
B	C	D	E	F	G	A
C	D	E	F	G	A	B
D	E	F	G	A	B	C
E	F	G	A	B	C	D
F	G	A	B	C	D	E
G	A	B	C	D	E	F

 8×8

A	B	C	D	E	F	G	H
B	C	D	E	F	G	H	A
C	D	E	F	G	H	A	B
D	E	F	G	H	A	B	C
E	F	G	H	A	B	C	D
F	G	H	A	B	C	D	E
G	H	A	B	C	D	E	F
H	A	B	C	D	E	F	G

 9×9

A	B	C	D	E	F	G	H	I
B	C	D	E	F	G	H	I	A
C	D	E	F	G	H	I	A	B
D	E	F	G	H	I	A	B	C
E	F	G	H	I	A	B	C	D
F	G	H	I	A	B	C	D	E
G	H	I	A	B	C	D	E	F
H	I	A	B	C	D	E	F	G
I	A	B	C	D	E	F	G	H

 10×10

A	B	C	D	E	F	G	H	I	J
B	C	D	E	F	G	H	I	J	A
C	D	E	F	G	H	I	J	A	B
D	E	F	G	H	I	J	A	B	C
E	F	G	H	I	J	A	B	C	D
F	G	H	I	J	A	B	C	D	E
G	H	I	J	A	B	C	D	E	F
H	I	J	A	B	C	D	E	F	G
I	J	A	B	C	D	E	F	G	H
J	A	B	C	D	E	F	G	H	I

 11×11

A	B	C	D	E	F	G	H	I	J	K
B	C	D	E	F	G	H	I	J	K	A
C	D	E	F	G	H	I	J	K	A	B
D	E	F	G	H	I	J	K	A	B	C
E	F	G	H	I	J	K	A	B	C	D
F	G	H	I	J	K	A	B	C	D	E
G	H	I	J	K	A	B	C	D	E	F
H	I	J	K	A	B	C	D	E	F	G
I	J	K	A	B	C	D	E	F	G	H
J	K	A	B	C	D	E	F	G	H	I
K	A	B	C	D	E	F	G	H	I	J

Plan 4.1 (Continued)

Selected latin squares

 12×12

A	B	C	D	E	F	G	H	I	J	K	L
B	C	D	E	F	G	H	I	J	K	L	A
C	D	E	F	G	H	I	J	K	L	A	B
D	E	F	G	H	I	J	K	L	A	B	C
E	F	G	H	I	J	K	L	A	B	C	D
F	G	H	I	J	K	L	A	B	C	D	E
G	H	I	J	K	L	A	B	C	D	E	F
H	I	J	K	L	A	B	C	D	E	F	G
I	J	K	L	A	B	C	D	E	F	G	H
J	K	L	A	B	C	D	E	F	G	H	I
K	L	A	B	C	D	E	F	G	H	I	J
L	A	B	C	D	E	F	G	H	I	J	K

Plan 4.2

Graeco-latin squares

 3×3

A ₁	B ₃	C ₂
B ₂	C ₁	A ₃
C ₃	A ₂	B ₁

 4×4

A ₁	B ₃	C ₄	D ₂
B ₂	A ₄	D ₃	C ₁
C ₃	D ₁	A ₂	B ₄
D ₄	C ₂	B ₁	A ₃

 5×5

A ₁	B ₃	C ₅	D ₂	E ₄
B ₂	C ₄	D ₁	E ₃	A ₅
C ₃	D ₅	E ₂	A ₄	B ₁
D ₄	E ₁	A ₃	B ₅	C ₂
E ₅	A ₂	B ₄	C ₁	D ₃

 7×7

A ₁	B ₅	C ₂	D ₆	E ₃	F ₇	G ₄
B ₂	C ₆	D ₃	E ₇	F ₄	G ₁	A ₅
C ₃	D ₇	E ₄	F ₁	G ₅	A ₂	B ₆
D ₄	E ₁	F ₅	G ₂	A ₆	B ₃	C ₇
E ₅	F ₂	G ₆	A ₃	B ₇	C ₄	D ₁
F ₆	G ₃	A ₇	B ₄	C ₁	D ₅	E ₂
G ₇	A ₄	B ₁	C ₅	D ₂	E ₆	F ₃

 8×8

A ₁	B ₅	C ₂	D ₃	E ₇	F ₄	G ₈	H ₆
B ₂	A ₈	G ₁	F ₇	H ₃	D ₆	C ₅	E ₄
C ₃	G ₄	A ₇	E ₁	D ₂	H ₅	B ₆	F ₈
D ₄	F ₃	E ₆	A ₅	C ₈	B ₁	H ₇	G ₂
E ₅	H ₁	D ₈	C ₄	A ₆	G ₃	F ₂	B ₇
F ₆	D ₇	H ₄	B ₈	G ₅	A ₂	E ₃	C ₁
G ₇	C ₆	B ₃	H ₂	F ₁	E ₈	A ₄	D ₅
H ₈	E ₂	F ₅	G ₆	B ₄	C ₇	D ₁	A ₃

 9×9

A ₁	B ₃	C ₂	D ₇	E ₉	F ₈	G ₄	H ₆	I ₅
B ₂	C ₁	A ₃	E ₈	F ₇	D ₉	H ₅	I ₄	G ₆
C ₃	A ₂	B ₁	F ₉	D ₈	E ₇	I ₆	G ₅	H ₄
D ₄	E ₆	F ₅	G ₁	H ₃	I ₂	A ₇	B ₉	C ₈
E ₅	F ₄	D ₆	H ₂	I ₁	G ₃	B ₈	C ₇	A ₉
F ₆	D ₅	E ₄	I ₃	G ₂	H ₁	C ₉	A ₈	B ₇
G ₇	H ₉	I ₈	A ₄	B ₆	C ₅	D ₁	E ₃	F ₂
H ₈	I ₇	G ₉	B ₅	C ₄	A ₆	E ₂	F ₁	D ₃
I ₉	G ₈	H ₇	C ₆	A ₅	B ₄	F ₃	D ₂	E ₁

Plan 4.2 (Continued)

Graeco-latin squares

 11×11

A ₁	B ₇	C ₂	D ₈	E ₃	F ₉	G ₄	H ₁₀	I ₅	J ₁₁	K ₆
B ₂	C ₈	D ₃	E ₉	F ₄	G ₁₀	H ₅	I ₁₁	J ₆	K ₁	A ₇
C ₃	D ₉	E ₄	F ₁₀	G ₅	H ₁₁	I ₆	J ₁	K ₇	A ₂	B ₈
D ₄	E ₁₀	F ₅	G ₁₁	H ₆	I ₁	J ₇	K ₂	A ₃	B ₃	C ₉
E ₅	F ₁₁	G ₆	H ₁	I ₇	J ₂	K ₃	A ₄	B ₉	C ₄	D ₁₀
F ₆	G ₁	H ₇	I ₂	J ₈	K ₄	A ₉	B ₄	C ₁₀	D ₅	E ₁₁
G ₇	H ₂	I ₃	J ₃	K ₉	A ₄	B ₁₀	C ₅	D ₁₁	E ₆	F ₁
H ₈	I ₃	J ₉	K ₄	A ₁₀	B ₅	C ₁₁	D ₆	E ₁	F ₇	G ₂
I ₉	J ₄	K ₁₀	A ₅	B ₁₁	C ₆	D ₁	E ₇	F ₂	G ₈	H ₃
J ₁₀	K ₅	A ₁₁	B ₆	C ₁	D ₇	E ₂	F ₈	G ₃	H ₉	I ₄
K ₁₁	A ₆	B ₁	C ₇	D ₂	E ₈	F ₃	G ₉	H ₄	I ₁₀	J ₅

 12×12

A ₁	B ₁₂	C ₆	D ₇	I ₅	J ₄	K ₁₀	L ₁₁	E ₉	F ₈	G ₂	H ₃
B ₂	A ₁₁	D ₅	C ₈	J ₆	I ₃	L ₉	K ₁₂	F ₁₀	E ₇	H ₁	G ₄
C ₃	D ₁₀	A ₈	B ₅	K ₇	L ₂	I ₁₂	J ₉	G ₁₁	H ₆	E ₄	F ₁
D ₄	C ₉	B ₇	A ₆	L ₈	K ₁	J ₁₁	I ₁₀	H ₁₂	G ₅	F ₃	E ₂
E ₅	F ₄	G ₁₀	H ₁₁	A ₉	B ₈	C ₂	D ₃	I ₁	J ₁₂	K ₆	L ₇
F ₆	E ₃	H ₉	G ₁₂	B ₁₀	A ₇	D ₁	C ₄	J ₂	I ₁₁	L ₅	K ₈
G ₇	H ₂	E ₁₂	F ₉	C ₁₁	D ₆	A ₄	B ₁	K ₃	L ₁₀	I ₈	J ₅
H ₈	G ₁	F ₁₁	E ₁₀	D ₁₂	C ₅	B ₃	A ₂	L ₄	K ₉	J ₇	I ₆
I ₉	J ₃	K ₂	L ₃	E ₁	F ₁₂	G ₆	H ₇	A ₅	B ₄	C ₁₀	D ₁₁
J ₁₀	I ₇	L ₁	K ₄	F ₂	E ₁₁	H ₅	G ₈	B ₆	A ₃	D ₉	C ₁₂
K ₁₁	L ₆	I ₄	J ₁	G ₃	H ₁₀	E ₈	F ₅	C ₇	D ₂	A ₁₂	B ₉
L ₁₂	K ₅	J ₃	I ₂	H ₄	G ₉	F ₇	E ₆	D ₈	C ₁	B ₁₁	A ₁₀

CHAPTER 5

FACTORIAL EXPERIMENTS

5.1 Description

5.11 A 2^2 Factorial Experiment. In a factorial experiment the effects of a number of different factors are investigated simultaneously. The treatments consist of all combinations that can be formed from the different factors. To illustrate the simplest case, consider an experiment on sugar beet with 2 factors. These were nitrogen, none (n_0) versus 3 cwt. sulphate of ammonia per acre (n_1), and depth of winter ploughing (7 in. versus 11 in.). Ploughing took place in late January, the nitrogen was applied in late April, and the seed was sown early in May. Since both factors occur at 2 levels or variations, the experiment is described as a 2×2 factorial experiment. The 4 treatment combinations are shown below, with the mean yields of sugar per acre (cwt.) underneath.

Treatment combinations and yields of sugar (cwt. per acre)

1	2	3	4
(n_0 , 7 in.)	(n_1 , 7 in.)	(n_0 , 11 in.)	(n_1 , 11 in.)
40.9	47.8	42.4	50.2

The yields may be placed in the following 2×2 table.

Depth	Nitrogen		Mean	Response to n_1
	n_0	n_1		
7 in.	40.9	47.8	44.4	+6.9
11 in.	42.4	50.2	46.3	+7.8
Mean	41.6	49.0	45.3	
11 in. minus 7 in.	+1.5	+2.4		

The results might be summarized as follows. Considering the effect of nitrogen, we might report that the application of nitrogen increased yields by 6.9 cwt. with shallow ploughing and by 7.8 cwt. with deep ploughing. These figures are called the *simple* effects of nitrogen. They

represent the type of information that would be wanted, for instance, in giving advice to a farmer who always used shallow ploughing but was doubtful whether to apply nitrogen. For the simple effects of depth of ploughing, we might report that 11 in. ploughing was superior to 7 in. by 1.5 cwt. in the absence of nitrogen and by 2.4 cwt. when nitrogen was applied.

There is another way of looking at the results. It sometimes happens that the effects of the factors are *independent*. By this we mean that the response to nitrogen is the same whether ploughing is shallow or deep, and that the difference between the effects of deep and shallow ploughing is the same whether nitrogen is present or not. In this event the two simple effects of nitrogen, 6.9 cwt. and 7.8 cwt., are estimates of the same quantity and differ only by experimental errors. On this supposition we would naturally average the two figures in order to estimate the response to nitrogen. The average, 7.4 cwt., is called the *main* effect of nitrogen. It can be derived alternatively as the difference between the two column means in the table, 41.6 and 49.0. Similarly the main effect of depth of ploughing (11 in. minus 7 in.) is the average of 1.5 cwt. and 2.4 cwt., or 1.9 cwt. Note that a main effect, being an average of the simple effects, is more precisely estimated than the latter. In this experiment the standard error of a main effect is $1/\sqrt{2}$ times that of a simple effect.

Consequently, if we are sure that the factors operate independently, the summary that was given above in terms of simple effects may be replaced by another that is both more concise and more accurate. This might read as follows. "The application of nitrogen increased the yield of sugar by 7.4 cwt., while 11 in. ploughing increased the yield by 1.9 cwt. as compared with 7 in. ploughing." It is worth repeating that when the factors are independent the figure 7.4 cwt. is the best estimate not only of the *average* response to nitrogen, but also of the response on plots ploughed to 7 in. and of that on plots ploughed to 11 in. In other words, the whole of the information in the experiment is contained in the main effects.

The question arises: How do we know whether the factors are independent? Frequently the answer is suggested by knowledge of the processes by which the factors produce their effects. In the present case an agronomist might reason that deep ploughing should enable the plant to develop a more vigorous root system. With this the plant should be able to utilize more effectively any added nutrient such as nitrogen. Thus he might predict that the response to nitrogen would be greater with deep than with shallow ploughing, though he probably would not expect it to be much greater. In short, he would predict that the two factors would not be quite independent in their effects.

In addition to the information that may be available from such reasoning, a factorial experiment itself provides a test of the assumption of independence. For, if the depth of ploughing does affect the response to nitrogen, the difference between 7.8 cwt. (the response to nitrogen with deep ploughing) and 6.9 cwt. (the response with shallow ploughing) is an estimate of this effect. The difference, 0.9 cwt., can be tested in the usual way by a *t*-test. If it proves significant, the assumption of independence is rejected by the data. The difference (sometimes divided by a numerical factor) is called the *interaction* between nitrogen and depth of ploughing.

Interchanging the roles of the two factors, we may also consider whether the superiority of deep over shallow ploughing is affected by the presence of nitrogen. To measure the interaction in this case, we subtract 1.5 cwt. (superiority of deep ploughing when no nitrogen is added) from 2.4 cwt. (superiority when nitrogen is added). The difference is again 0.9 cwt. It is easy to see that this equality always holds with a 2×2 experiment. Each difference is equal to the sum of the observations in one diagonal of the 2×2 table, minus the corresponding sum for the other diagonal. Such interactions are called *two-factor*, or *first-order*, interactions.

5.12 Advantages of Factorial Experimentation When Factors Are Independent. The advantages of factorial experimentation naturally depend on the purpose of the experiment. We suppose for the present that the purpose is to investigate the effects of each factor over some pre-assigned range that is covered by the levels of that factor which are used in the experiment. In other words the object is to obtain a broad picture of the effects of the factors rather than to find, say, the combination of the levels of the factors that give a maximum response. One procedure is to conduct separate experiments each of which deals only with a single factor. Another is to include all factors simultaneously by means of a factorial experiment.

If all factors are independent in their effects, the factorial approach will result in a considerable saving of the time and material devoted to the experiments. The saving follows from two facts. First, as we have seen, when factors are independent all the simple effects of a factor are equal to its main effect, so that main effects are the only quantities needed to describe fully the consequences of variations in the factor. Secondly, in a factorial experiment, each main effect is estimated with the same precision as if the whole experiment had been devoted to that factor alone. Thus, in the preceding example, half the plots receive nitrogen and half do not. Consequently, the main effect of nitrogen is estimated

just as precisely as it would be in a simple experiment of the same size devoted to nitrogen alone. The same result holds for the effect of depth of ploughing. The two single-factor experiments would require *twice* the total number of plots in order to equal the precision obtained by the factorial experiment. If there were n factors, all at two levels and all independent, the single-factor approach would necessitate n times as much experimental material as a factorial arrangement of equal precision. The gain from factorial arrangements in this case is very substantial.

Practical considerations may diminish this gain. The experimenter frequently lacks the resources to conduct a large experiment with many treatments, and must proceed with only one or two factors at a time. Further, it has been pointed out previously that, as the number of treatment combinations in an experiment is enlarged, the standard error per unit increases. This standard error is therefore likely to be higher for a large factorial experiment than for a comparable single-factor experiment. This increase in standard errors can usually be kept small by the device known as confounding, described in chapter 6.

5.13 Factorial Experimentation When Factors Are Not Independent.

We assume that the purpose is still to investigate each factor over the range represented by its levels. When factors are not independent, the simple effects of a factor vary according to the particular combination of the other factors with which these are produced. In this case the single-factor approach is likely to provide only a number of disconnected pieces of information that cannot easily be put together. In order to conduct an experiment on a single factor A , some decision must be made about the levels of other factors B , C , D , say, that are to be used in the experiment (e.g., whether all plots should be ploughed 7 in., 9 in., or 11 in. deep in an experiment on nitrogen). The experiment reveals the effects of A for this particular combination of B , C , and D , but no information is provided for predicting the effects of A with any other combination of B , C , and D . With a factorial approach, on the other hand, the effects of A are examined for every combination of B , C , and D that is included in the experiment. Thus a great deal of information is accumulated both about the effects of the factors and about their inter-relationships.

In this connection, Fisher (5.1) has pointed out that it is sometimes advisable to introduce into an experiment an extra factor that is not itself of interest, in order that the experiment may form the basis for sounder recommendations about the other factors. In agricultural experimentation in Britain, farmyard manure has served as a subsidiary factor of this kind. Any recommendations made to farmers about other

factors will be put to the test in some fields where the farmer has applied manure and in others where he has not, so that it is well to investigate the other factors in both the presence and the absence of manure.

5.14 Summary Comments. To summarize, the following are some instances where factorial experimentation may be suitable:

1. In exploratory work where the object is to determine quickly the effects of each of a number of factors over a specified range.

2. In investigations of the interactions among the effects of several factors. From their nature, interactions cannot be studied without testing *some* of the combinations formed from the different factors. Frequently, information is most quickly obtained by testing all combinations.

3. In experiments designed to lead to recommendations that must apply over a wide range of conditions. Subsidiary factors may be brought into an experiment so as to test the principal factors under a variety of conditions similar to those that will be encountered in the population to which recommendations are to apply.

On the other hand, if considerable information has accumulated, or if the object of the investigation is specialized, it may be more profitable to conduct intensive work on a single factor or on a few combinations of factors. For instance, some investigations are directed towards finding the combination of the levels of the factors that will produce a maximum response. An interesting discussion of procedures for this purpose has been given by Friedman and Savage (5.2). They consider the case, common in industrial experimentation, where the effect of any specified treatment combination can be determined quickly. Thus the treatment combinations to be tested can be decided as the experiment proceeds, in the light of knowledge gained from combinations that have already been tested. They propose first a single-factor experiment to determine the optimum level a_1 of A for fixed levels b , c , and d , say, of the other factors. This is followed by a single-factor experiment to determine the optimum level b_1 of B for fixed values a_1 , c , and d . Similarly, C is tested at a_1 , b_1 , d , and D at a_1 , b_1 , c_1 . After the completion of the first "round," the whole process is repeated until the maximum appears to have been reached. The authors show that the maximum will usually be reached more quickly than with a complete factorial, since the plan is designed to concentrate on combinations that are near the maximum. The plan would not be feasible for agricultural field experiments, where in general an experiment can be changed only once a year.

Experimenters sometimes find the results of factorial experiments

difficult to interpret, because they appear to present a bewildering variety of treatment comparisons. It is true that the competent summary of a large factorial experiment demands an orderly procedure and often takes considerable time. If the factors are for the most part independent, the method of analysis by means of main effects and interactions (to be illustrated later) will reduce the data to manageable proportions. If the numerous factors interact in a puzzling manner, prolonged study of the results and further experimentation may be needed before the facts are mastered. The trouble in this case is that the phenomena are complex, not that the experimentation is faulty.

5.2 Calculation of Main Effects and Interactions

5.21 Notation for the 2ⁿ Series. The object of section 5.2 is to explain how main effects and interactions are calculated, and how they are represented in the analysis of variance of the results. We begin with the 2ⁿ series, where each factor occurs at only two levels.

For this system the notation used is similar to that of Yates (5.3). Letters A, B, C, \dots denote the factors. The letters a, b, c, \dots denote one of the two levels at which the corresponding factor occurs; for purposes of clarity this level will be called the second level. The first level is signified by absence of the corresponding letter. Thus the treatment combination bd , in a 2⁴ factorial experiment, means the treatment combination which contains the first levels of factors A and C , and the second levels of factors B and D . The treatment combination which consists of the first level of *all* factors is denoted by the symbol (1).

The symbol (ab) will denote the *mean* of all observations which receive the treatment combination ab . The letters A, B , and AB , when they refer to numbers, will represent, respectively, the main effects of A and B and the A by B interaction.

5.22 The 2² Factorial Experiment. As already shown in section 5.11, we have

$$A = \frac{1}{2}[(ab) - (b) + (a) - (1)]$$

$$B = \frac{1}{2}[(ab) + (b) - (a) - (1)]$$

Yates (5.3) introduces the same multiplier $\frac{1}{2}$ for the interaction, which he defines as

$$AB = \frac{1}{2}[(ab) - (b) - (a) + (1)]$$

These quantities and the general mean M are shown in terms of the means for the treatment combinations in table 5.1.

TABLE 5.1 MAIN EFFECTS AND INTERACTIONS EXPRESSED IN TERMS OF INDIVIDUAL TREATMENT MEANS: 2^2 FACTORIAL

Factorial effect	Treatment combination				Divisor
	(1)	(a)	(b)	(ab)	
<i>M</i>	+	+	+	+	4
<i>A</i>	-	+	-	+	2
<i>B</i>	-	-	+	+	2
<i>AB</i>	+	-	-	+	2

The rows of the table express the factorial effects in terms of the original means. If the equations represented by the table are solved for (ab) , (b) , etc., in terms of M , A , etc., it will be found that the columns of the table enable us to express the original means in terms of the factorial effects. For example, from the column for (ab) ,

$$(ab) = M + \frac{1}{2}[A + B + AB]$$

The only point to remember is that the factor $\frac{1}{2}$ occurs with all terms except M . From these results, simple effects may be calculated from factorial effects. Thus

$$(a) - (1) = \text{simple effect of } A \text{ when } B \text{ is at the first level} = A - AB$$

$$(ab) - (b) = \text{simple effect of } A \text{ when } B \text{ is at the second level} = A + AB$$

These relations are useful when an experiment has been summarized in terms of the factorial effects, and it is later desired to estimate some of the simple effects. From the above example it may be noted that the quantity AB measures the error that is committed in estimating the simple effects of A if the two factors are erroneously assumed to be independent.

The functions A , B , and AB satisfy the conditions for an orthogonal set of functions (section 3.42). Consequently the squares of the factorial effects, when suitably multiplied, divide the treatments s.s. into three single components, each with 1 d.f. In practice these components will usually be calculated from the treatment totals $[ab]$, $[b]$, etc., rather than from means. If we define factorial effect *totals* as illustrated below,

$$[A] = [ab] - [b] + [a] - [1]$$

then the sum of squares for A in the analysis of variance is $[A]^2/4r$, where r is the number of replicates. Corresponding formulae hold for the other factorial effects.

5.23 The 2³ Factorial Experiment. In this case there are *eight* treatment combinations: (1), a , b , c , ab , ac , bc , and abc . The simple effect of A is determined for each of *four* combinations of the other factors: (1), b , c , and bc . As before, the main effect of A is defined to be the average of these four simple effects.

$$A = \frac{1}{4}[(abc) - (bc) + (ab) - (b) + (ac) - (c) + (a) - (1)]$$

Similar definitions hold for B and C .

The interaction of A with B is now measured separately at each of the two levels of C .

$$AB \text{ (C at second level)} = \frac{1}{2}[(abc) - (bc) - (ac) + (c)]$$

$$AB \text{ (C at first level)} = \frac{1}{2}[(ab) - (b) - (a) + (1)]$$

As would be expected, the quantity AB is taken as the average of these two effects. Thus

$$AB = \frac{1}{4}[(abc) - (bc) - (ac) + (c) + (ab) - (b) - (a) + (1)]$$

There are two other first-order interactions, AC and BC , which are defined similarly.

In addition, we encounter a new interaction. Separate estimates were given above for AB at each of the two levels of C . The difference between these two estimates measures the effect of C on the AB interaction. This difference, with the conventional factor $\frac{1}{4}$, is

$$\frac{1}{4}[(abc) - (bc) - (ac) + (c) - (ab) + (b) + (a) - (1)]$$

and may be called the interaction of AB with C . If the algebra is carried out, it will be found that the same expression measures the interaction of AC with B , and that of BC with A . Hence the quantity is called the ABC interaction. It is a *three-factor*, or *second-order*, interaction.

Three-factor interactions are more difficult to understand than two-factor interactions. Fortunately, in practice three-factor interactions are often small relative to main effects and two-factor interactions; and quite frequently they can be neglected for the purposes to which the results are to be put. Occasionally, cases arise where they are important. It might happen, for instance, that factor A does not exert any influence unless factors B and C are present in the combination (bc) . In this event the interaction ABC is as large as the main effect of A or the interactions AB and AC . The same effect may occur in less extreme cases, where the

combination bc is particularly favorable to the response to factor A . If three-factor interactions are found to be substantial, a careful scrutiny of the simple effects is usually helpful in interpretation.

The expressions for the factorial effects in terms of the treatment means are summarized in table 5.2. The rows of the table give the

TABLE 5.2 MAIN EFFECTS AND INTERACTIONS EXPRESSED IN TERMS OF INDIVIDUAL TREATMENT MEANS: 2^3 FACTORIAL

Factorial effect	Treatment combination								Divisor
	(1)	(a)	(b)	(ab)	(c)	(ac)	(bc)	(abc)	
M	+	+	+	+	+	+	+	+	8
A	-	+	-	+	-	+	-	+	4
B	-	-	+	+	-	-	+	+	4
C	-	-	-	-	+	+	+	+	4
AB	+	-	-	+	+	-	-	+	4
AC	+	-	+	-	-	+	-	+	4
BC	+	+	-	-	-	-	+	+	4
ABC	-	+	+	-	+	-	-	+	4

factorial effects in terms of the treatment means, while the columns give the treatment means in terms of the factorial effects. For example,

$$(a) = M + \frac{1}{2}[A - B - C - AB - AC + BC + ABC]$$

$$(a) - (1) = A - AB - AC + ABC$$

$$(abc) - (1) = A + B + C + ABC$$

As before, the factor $\frac{1}{2}$ appears with all terms except M in the expression for a treatment mean. The difference between two treatment means, of which two examples are given above, is easily found by noting the signs in the two columns in question.

As the reader may verify, the 7 factorial effects are mutually orthogonal, and each is orthogonal to M . If r is the number of replicates, each factorial effect has the same variance, $\frac{8\sigma^2}{4^2r}$, or $\frac{\sigma^2}{2r}$. The contribution of any effect to the sum of squares for treatments is $[]^2/8r$, where $[]$ denotes the factorial effect total; e.g.,

$$[A] = -[1] + [a] - [b] + [ab] - [c] + [ac] - [bc] + [abc]$$

5.24 The 2^n Factorial Experiment. A few formulae will be given which apply to any number of factors, all at two levels. With n factors, there are n main effects, $n(n-1)/2$ two-factor interactions, $n(n-1)(n-2)/6$ three-factor interactions, and so on. The successive numbers are the coefficients in the expansion of $(1+1)^n$, omitting the first coefficient, unity.

Two equivalent methods for writing out any factorial effect in terms of the original treatment means are available. Each is simple to remember, though the actual writing may take some time in a large experiment. They will be illustrated by finding the $BCDE$ interaction in a 2^6 experiment, with factors A, B, C, D, E , and F .

Rule 1. Evens versus odds. In every factorial effect, half the treatment combinations receive a $+$ sign and half a $-$ sign. Those which receive a $+$ sign are those which contain an *even* number of the letters that appear in the factorial effect. In the $BCDE$ interaction 4 letters, b, c, d , and e , appear. There are 8 combinations that contain an even number of letters: (1), bc, bd, be, cd, ce, de , and $bcde$. Each of these can be combined with any one of 4 combinations of the remaining letters a and f , namely, (1), a, f , and af . Hence, the 32 terms that appear with a $+$ sign are as follows.

(1)	a	f	af
bc	abc	bcf	$abcf$
bd	abd	bdf	$abdf$
be	abe	bef	$abef$
cd	acd	cdf	$acdf$
ce	ace	cef	$acef$
de	ade	def	$adef$
$bcde$	$abcde$	$bcdef$	$abcdef$

The 32 combinations with a $-$ sign are those which have an *odd* number of the letters b, c, d , or e . In detail, these are

b	ab	bf	abf
c	ac	cf	acf
d	ad	df	adf
e	ae	ef	aef
bcd	$abcd$	$bcdf$	$abcdf$
bce	$abce$	$bcef$	$abcef$
bde	$abde$	$bdef$	$abdef$
cde	$acde$	$cdef$	$acdef$

When calculated from the treatment *means*, this difference between the sums for the two groups of 32 combinations is divided by 2^{n-1} to give the $BCDE$ effect.

Rule 2. Algebraic. In this rule the *BCDE* interaction is expressed formally as

$$BCDE \approx \frac{1}{2^{n-1}} (a+1)(b-1)(c-1)(d-1)(e-1)(f+1)$$

Note that $-$ signs appear with the factors that enter into the interaction and $+$ signs with those that do not. If this expression is expanded algebraically, it gives the interaction as a linear function of the treatment means.

As in the 2^2 and 2^3 cases, all factorial effects are orthogonal to one another and to the mean. The contribution of any effect to the sum of squares in the analysis of variance is $[]^2/2^n r$, where $[]$ denotes the effect total.

5.25 Factors at More than Two Levels: a 4×2 Factorial. Various notations are used. For example, the 3 levels of a factor *A* may be denoted by a_0, a_1, a_2 , or by a_1, a_2, a_3 , or simply by the numbers 0, 1, 2. For illustration, we give below the treatment totals in a 4×2 experiment on sugar cane. The treatments were 4 levels of dressing of potash, k_0, k_1, k_2 , and k_3 , in arithmetic progression, and 2 levels of phosphate, p_0 and p_1 . There were 5 replicates in randomized blocks.

TABLE 5.3 TOTAL YIELDS OF SUGAR (TONS PER ACRE)

	k_0	k_1	k_2	k_3	Total
p_0	180	248	277	285	990
p_1	251	307	342	346	1246
Totals	431	555	619	631	2236

The main effect of phosphate (*P*), which occurs at only 2 levels, is of course derived from the comparison of the 2 marginal totals, 990 and 1246. For the main effect of potash (*K*) there are 4 marginal totals which may be compared. It will be recalled (section 3.42) that 3 independent comparisons may be made amongst 4 totals, and that an infinite number of such sets of 3 may be chosen. Thus the main effect of *K* comprises 3 independent comparisons.

With the 2^n system, we were able to define every main effect as a specific linear combination of the treatment means. In the present case we could select a particular set of 3 independent comparisons, each of which would be a specific linear combination of the treatment means. These could be defined as the "components" of the main effect of *K*. However, the particular set that is most useful for the interpretation of the results

will change from experiment to experiment, so that a formal definition of this type would be of limited utility. The experimenter should use whichever set appears most relevant.

Now consider the interaction between P and K . From the differences between the two rows in table 5.3, the effect of P is estimated separately at each level of K , the estimates being

	k_0	k_1	k_2	k_3
$(p_1 - p_0)$	71	59	65	61

Any comparison among these 4 figures is a measure of the effect of K on the response to P , and therefore is a component of the interaction between P and K . Consequently the interaction between P and K consists of 3 independent comparisons, and is said to have 3 d.f. As with the main effect of K , the particular set of 3 components that will be of interest varies with the type of experiment.

We may also wish to consider the interaction as the effect of P on the response to K . As we have seen, the response to K has 3 components. Since there are 4 increasing levels of potash dressing, we might choose as components the linear, quadratic, and cubic components of the response curve. Apart from a divisor, the linear component is

$$-3(180) - 1(248) + 1(277) + 3(285) = 344$$

at the lower level of P and

$$-3(251) - 1(307) + 1(342) + 3(346) = 320$$

at the higher level of P . The difference between these 2 figures, -24 , estimates the effect of P on the linear response to K , and is a part of the interaction between K and P . The other two components are the effects of P on the quadratic and cubic responses to K .

In a 2^n system, the interaction between P and K is identical with the interaction between K and P . In the more general case the corresponding result is that any component of the interaction between K and P can be derived from the components of the interaction between P and K . For example, the effect of P on the linear response to K can be written

$$-24 = -3(71) - 1(59) + 1(65) + 3(61)$$

so that it is a linear function of the responses to P at the 4 levels of K . In this more general sense, the two interactions are still equivalent.

We will consider the analysis of variance of the 8 treatment totals in some detail. With 5 replicates, the total s.s., on a single-plot basis, is

$$\frac{(180)^2 + (251)^2 + (248)^2 + \dots + (346)^2}{5} - \frac{(2236)^2}{40} = 4165.2$$

The 7 d.f. subdivide into 1 for the main effect of P , 3 for that of K , and 3 for the PK interaction. In practice the first two terms are calculated directly and the interaction obtained by subtraction. The computations are

$$P: \frac{(1246 - 990)^2}{40} = 1638.4$$

$$K: \frac{(431)^2 + (555)^2 + (619)^2 + (631)^2}{10} - \frac{(2236)^2}{40} = 2518.4$$

TABLE 5.4 PRELIMINARY ANALYSIS OF VARIANCE OF DATA IN TABLE 5.3

	d.f.	s.s.	m.s.
P	1	1638.4	1638.4
K	3	2518.4	839.5
PK	3	8.4	2.8
	—	—	—
Total	7	4165.2	

As is common in agricultural experiments, the interaction m.s. is small compared with those for the main effects (in fact it is below the mean square for error).

As an exercise we will divide the K main effect and its interaction with P into linear, quadratic, and cubic components. The first step is to calculate the totals for these effects separately at each level of P . The results are shown below, the multipliers for the 4 levels of K being shown in parentheses. These multipliers are obtained from a table of orthogonal polynomials (5.9).

	K_l (-3, -1, +1, +3)	K_q (+1, -1, -1, +1)	K_c (-1, +3, -3, +1)
p_0	+344	- 60	+18
p_1	+320	- 52	-10
	—	—	—
Sum ($p_1 + p_0$)	+664	-112	+ 8
Diff. ($p_1 - p_0$)	- 24	+ 8	-28

From the *sum* line, we obtain the 3 components of the K main effect,

$$K_l = \frac{1}{200}(664)^2 = 2204.5, \quad K_q = \frac{1}{40}(112)^2 = 313.6$$

$$K_c = \frac{1}{200}(8^2) = 0.3$$

Similarly, from the *difference* line, the components of the PK interaction are

$$PK_l = \frac{1}{200}(24)^2 = 2.9, \quad PK_q = \frac{1}{40}(8)^2 = 1.6$$

$$PK_c = \frac{1}{200}(28)^2 = 3.9$$

The divisors are found from the usual rule for linear functions (section 3.42). The three K components add to 2518.4, the total s.s. for K , while the PK components add to 8.4, the sum of squares for PK . The detailed analysis of variance is shown in table 5.5.

TABLE 5.5 MORE DETAILED ANALYSIS OF VARIANCE OF THE DATA
IN TABLE 5.3

	d.f.	s.s. or m.s.
P	1	1638.4
K_l	1	2204.5
K_q	1	313.6
K_c	1	0.3
PK_l	1	2.9
PK_q	1	1.6
PK_c	1	3.9

Since every component has 1 d.f., the mean squares are the same as the sums of squares. The error m.s. is 16.35, with 28 d.f. The quadratic component of K is significant, indicating a falling off in the response at the higher levels of application. Neither the cubic component nor any of the interactions approaches significance.

5.26 A 3² Factorial Experiment. As a second example, the treatment totals are shown in table 5.6 for an experiment with 3 levels of nitrogen fertilizer and 3 of phosphate fertilizer. The data are the numbers of lettuce plants that emerged from the ground and are totals over 12 plots

TABLE 5.6 NUMBERS OF LETTUCE PLANTS EMERGING *

	n_0	n_1	n_2	Totals
p_0	449 ($A\alpha$)	413 ($C\beta$)	326 ($B\gamma$)	1188
p_1	409 ($B\beta$)	358 ($A\gamma$)	291 ($C\alpha$)	1058
p_2	341 ($C\gamma$)	278 ($B\alpha$)	312 ($A\beta$)	931
Totals	1199	1049	929	3177

* The use of the latin and greek letters will be explained later.

each. Both nitrogen and phosphate appear to have had a deleterious effect on emergence (the subscript 2 denotes the largest application). The main effects of N and P both comprize two independent comparisons, and thus have 2 d.f. each. Since the amounts of N and P were in arithmetic progression, it would probably again be appropriate, as in the previous example, to choose the linear and quadratic components of the regression on amount of dressing as the individual components of the main effects.

The N_l component can be estimated separately at each level of P . This means that there are 2 d.f. which measure the effect of P on the linear response to N . These are a part of the NP interaction. Another 2 d.f. are supplied by the effect of P on the N_q component, so that the NP interaction contains 4 d.f.

From inspection of the margins of table 5.6, it is evident that the main effects of both N and P are approximately linear. Consequently, the most interesting single degree of freedom from the interactions is likely to be the interaction of N_l with P_l . In many agricultural experiments this type of interaction is the only one that approaches significance. It is worth while calculating this interaction separately in the analysis of variance. The other three components, N_lP_q , N_qP_l , and N_qP_q , will also be obtained. The initial computations appear in table 5.7.

TABLE 5.7 CALCULATION OF LINEAR AND QUADRATIC EFFECTS FOR THE ANALYSIS OF VARIANCE

	N_l (-1, 0, 1)	N_q (1, -2, 1)		P_l (-1, 0, 1)	P_q (1, -2, 1)
p_0	-123	-51	n_0	-108	-28
p_1	-118	-16	n_1	-135	-25
p_2	-29	+97	n_2	-14	+56
Sum	$-270(N_l)$	$+30(N_q)$	Sum	$-257(P_l)$	$+3(P_q)$
P_l	$+94(N_lP_l)$	$+148(N_qP_l)$	N_l	$+94(P_lN_l)$	$+84(P_qN_l)$
P_q	$+84(N_lP_q)$	$+78(N_qP_q)$	N_q	$+148(P_lN_q)$	$+78(P_qN_q)$

The left side of the table shows the N_l and N_q effects for each level of P , while the right side shows the P_l and P_q effects for each level of N . For instance, from table 5.6,

$$N_l p_0 = 326 - 449 = -123; \quad P_q n_0 = 449 - 2(409) + 341 = -28$$

The column sums give the individual components of the N and P main effects. Now consider the difference between the third and the first row. For the N_l column, this difference (+94) gives the linear effect of P on N : i.e., the N_lP_l interaction. From the P_l column, the difference gives the linear effect of N on P_l , or the P_lN_l interaction, which is exactly the same as the N_lP_l interaction. The other two columns provide the N_qP_l and the N_lP_q effects.

Finally, the sum of the first and third rows, minus twice the second row, leads to the components of interaction that contain a quadratic term. It will be seen that all four components can be obtained from either the left or the right half of the table, so that in practice only one half is required. By computing both halves we verify the symmetry of the components with respect to N and P .

The squares of these quantities, with appropriate divisors, will give an analysis of variance of the 8 d.f. amongst the treatment totals into 8 single components. Since an individual entry in table 5.6 is the total of 12 plots, the divisors may be verified to be as shown below.

	N_l or P_l	N_q or P_q	N_lP_l	N_lP_q	N_qP_l	N_qP_q
Divisor	72	216	48	144	144	432

For instance, the divisor for N_qP_q may be worked out as follows. The three N_q figures in table 5.7 (i.e., -51, -16, +97) each have divisor $12(1^2 + 2^2 + 1^2)$, or 72. Since the N_qP_q total is a linear function of these three figures, with coefficients 1, -2, and 1, this total has divisor $72(1^2 + 2^2 + 1^2)$, or 432. The analysis of variance is shown in table 5.8.

TABLE 5.8 SUBDIVISION OF THE TREATMENT S.S.

	d.f.	s.s. or m.s.
N_l	1	1012.50
N_q	1	4.17
P_l	1	917.35
P_q	1	0.04
N_lP_l	1	184.08
N_lP_q	1	49.00
N_qP_l	1	152.11
N_qP_q	1	14.08

The error m.s. in this experiment is about 59. The linear effects of both fertilizers are significant, with no indication of curvature. The N_lP_l effect is significant at the 10% level, but not at the 5% level.

In table 5.6, p. 135, latin and greek letters were superimposed so as to form a 3×3 graeco-latin square. This square leads to another method of calculating the sum of squares for the interactions (4 d.f.). Although the method is not likely to be of use for purposes of interpretation, it has formed the basis of some ingenious devices in the construction of designs. In the square, the column totals represent the main effects of N and the row totals those of P . Since the latin letter totals are orthogonal to rows and columns, it seems reasonable to suppose that they must represent two of the 4 components of the NP interaction. Similarly, the greek letter totals represent the remaining 2 components. These totals are shown below.

A	B	C	Total	α	β	γ	Total
1119	1013	1045	3177	1018	1134	1025	3177

Each figure is now a total of 36 plots. The sum of squares of deviations of the latin letter totals, divided by 36, is 164.22, and the corresponding figure for the greek letters is 235.06. These add to 399.28, which is the same as the total s.s. for the interactions in table 5.8, apart from rounding differences.

5.27 General Method of Analysis. Suppose that there are three factors, A , B , C , which occur at α , β , and γ levels respectively. The main effects have $(\alpha - 1)$, $(\beta - 1)$, and $(\gamma - 1)$ components or degrees of freedom respectively. Each component of the main effect of A can be estimated separately at each of the β levels of B . Thus each component of A contributes $(\beta - 1)$ degrees of freedom to the AB interaction. This means that the AB interaction contains a total of $(\alpha - 1)(\beta - 1)$ components or degrees of freedom. Similarly the AC interaction has $(\alpha - 1)(\gamma - 1)$ degrees of freedom, and the BC interaction has $(\beta - 1)(\gamma - 1)$ degrees of freedom.

To compute the sums of squares for the main effects and first-order interactions, we form two-way tables for *each pair of factors*. Consider the A by B two-way table. The total s.s. among cells has $(\alpha\beta - 1)$ degrees of freedom. From the marginal totals in the table we compute the sum of squares for the main effect of A , with $(\alpha - 1)$ degrees of freedom, and that for the main effect of B , with $(\beta - 1)$ degrees of freedom. By subtraction, the sum of squares for the AB interaction, with $(\alpha - 1)(\beta - 1)$ degrees of freedom, is obtained.

There remains the three-factor, or ABC , interaction. Now each of the $(\alpha - 1)(\beta - 1)$ components in the AB interaction is estimated separately at each level of C . It will therefore contribute $(\gamma - 1)$ degrees of freedom to the ABC interaction, so that the latter contains in all $(\alpha - 1)(\beta - 1)(\gamma - 1)$ degrees of freedom. The sum of squares for ABC is also obtained most easily by subtraction. From the total s.s. amongst treatments, with $(\alpha\beta\gamma - 1)$ degrees of freedom, subtract the sums of squares for A , B , C , AB , AC , and BC . The remainder will be the sum of squares for ABC .

It is hoped that the reader will find no difficulty in extending these methods to the case where there are more than three factors. In general, the sums of squares for main effects are calculated directly, and those for interactions are calculated by subtraction. To compute an $ABCD$ interaction, for instance, we require a four-way table for the four factors represented. From the sum of squares for this table we subtract the sum of squares for all main effects and two- and three-factor interactions among the factors in question. Calculations must be checked by re-computation. Of course, if the experimenter subdivides any interaction

into single components as in the previous section, a check is provided by this process.

In the general case it remains true that any component of a factorial effect is orthogonal to any component of any *other* factorial effect. Thus any component of the *BC* interaction is orthogonal to any component of the *A* main effect, or of the *BCD* interaction, etc. Two components of the *same* factorial effect may or may not be independent. For instance, if *A* occurs at three levels, the comparison ($a_2 - a_0$) is independent of the comparison ($a_2 - 2a_1 + a_0$), but is not independent of ($a_1 - a_0$).

The following selected references contain a discussion of factorial experiments with some worked examples.

- 5.3 YATES, F. The design and analysis of factorial experiments. *Imp. Bur. Soil Sci. Tech. Comm.* 35, 1937. This gives the most comprehensive account that is available, with numerous worked examples.
- 5.4 YATES, F. Complex experiments. *Jour. Roy. Stat. Soc. Suppl.* 2, 181-247, 1935. An earlier reference, with examples of 2^2 , 2^3 , 4×3 , and 3^3 factorials.
- 5.5 TIPPETT, L. H. C. *The methods of statistics*. Williams and Norgate, London, 2nd ed., 1937. The use of factorial designs in industrial experimentation is indicated. A worked example is given of the breakdown of the treatment s.s. in a $5 \times 4 \times 3 \times 2$ factorial in single replication.
- 5.6 LINDQUIST, E. F. *Statistical analysis in educational research*. Houghton Mifflin, Boston, 1940. The application of factorial designs in experiments on methods of teaching is described, with a worked analysis of variance for a 4×3 design completely randomized.
- 5.7 BLISS, C. I. Factorial design and covariance in the biological assay of vitamin D. *Jour. Amer. Stat. Assoc.* 35, 498-506, 1940. Worked example of a 3×2 design.

5.28 Interpretation of the Analysis: First Example. The separation of the treatment comparisons into main effects and interactions is a convenient and powerful method of analysis in cases where interactions are small relative to main effects. When interactions are large, this analysis must be supplemented by a detailed examination of the nature of the interactions. It may, in fact, be found that an analysis into main effects and interactions is not suited to the data at hand. There is sometimes a tendency to apply the factorial method of analysis mechanically without considering whether it is suitable or not, and also a tendency to rely too much on the initial analysis of variance alone when writing a summary of the results. Below two examples are presented where the initial analysis of variance is not very informative, and where the results can be summarized better in terms of simple rather than factorial effects.

The first experiment was conducted by the Wailuku Sugar Company. Three varieties of sugar cane were compared, in combination with three levels of nitrogen (150, 210, and 270 lb. *N* per acre respectively). The

crop was the second harvesting, or the first ratoon crop. In table 5.9 only the relevant part of the analysis of variance is shown: i.e., the sum of squares for the main effects of V and N and for the VN interaction, plus the error s.s. The data are in tons of cane per acre. The conclusions

TABLE 5.9 ANALYSIS OF VARIANCE OF A 3×3 SUGAR CANE EXPERIMENT

	d.f.	s.s.	m.s.
V	2	319.38	159.69 *
N	2	56.54	28.27
VN	4	559.79	139.95 *
Error	24	1053.84	43.91

from the analysis of variance are that the main effects of V and the VN interaction are both significant, but there is no sign of a main effect due to N . This statement tells little about the results of the experiment.

The treatment totals (over 4 replications) are shown below with their standard errors. Since the standard error per plot is $\sqrt{43.91} = 6.626$, the standard error for a treatment total is 13.3 as shown.

TABLE 5.10 TREATMENT TOTALS (TONS) IN A 3×3 SUGAR CANE EXPERIMENT (± 13.3)

	n_0	n_1	n_2	Total	s.e.
v_1	266.1	275.9	303.8	845.8	
v_2	245.8	250.2	281.7	777.7	± 23.0
v_3	274.4	258.1	231.6	764.1	
Total	786.3	784.2	817.1	2387.6	
s.e.		± 23.0			

Instead of having no effect, nitrogen has apparently given a steady increase in yields with the first two varieties, but a steady decrease with the third variety. Further, the significant main effects of varieties apply only to the average varietal yields over all 3 dressings of N , and not to yields with a particular rate of dressing. On the average v_1 gives a substantially higher yield than v_3 , but at the lowest level of N , v_3 is slightly above v_1 .

The subsequent analysis may be made either by means of t -tests applied to table 5.10 or by means of a further subdivision of the analysis of variance. It is of interest to examine the response to N for each variety separately. To test the linear responses by means of a t -test, we require the standard error for $(n_2 - n_0)$ as computed for each variety. This standard error is 13.3×1.414 , or 18.8, and since the 5% t -value for 24 d.f. is 2.064, the quantities $(n_2 - n_0)$ must attain the value 18.8×2.064 ,

or 38.8, in order to be significant at the 5% level. The actual values are

$$(n_2 - n_0)v_1 = +37.7; \quad (n_2 - n_0)v_2 = +35.9;$$

$$(n_2 - n_0)v_3 = -42.8$$

Thus neither of the increases with v_1 and v_2 quite reaches the significance level, though both are close to it. The decrease with v_3 is significant.

The test of linearity of the response curves is made by means of the quantity $(n_2 + n_0 - 2n_1)$, calculated for each variety. The standard error of this quantity is $13.3 \times \sqrt{6}$, or 32.6. The reader may verify from table 5.10 that none of the curvature terms even exceeds its standard error, so that there is no occasion to reject the hypothesis that the responses are linear.

It is not quite so clear what tests are appropriate for appraising the varietal differences. Since, however, the difference between v_1 and v_2 is very consistent at all levels of N , a t -test of the total difference is suggested. This difference is 68.1, and since the value required for significance is $23.0 \times 1.414 \times 2.064$, or 67.1, the superiority of v_1 over v_2 is just significant. Further, it is evident on inspection that v_3 does not differ significantly from the other varieties at either of the two lower levels of N . At the highest level, v_3 is significantly below both v_1 and v_2 .

When interactions are large, much care is required in the preparation of a statement that summarizes the results, and it is not easy to reach a form that is free from criticism. The following is a suggestion for this example.

"The increase in yield of cane to the highest dressing of N (270 lb. per acre) over the lowest dressing (150 lb. per acre) was 9.4 tons per acre with v_1 and 9.0 tons per acre with v_2 . Both increases just failed to be significant at the 5% level. With v_3 , on the other hand, the highest dressing of N decreased the yield significantly by 10.7 tons per acre as compared with the lowest dressing. For all three varieties the effects of N appeared to be proportional to the amount applied, within the range investigated in this experiment.

"Variety 1 gave a higher yield than variety 2 for all levels of N , the average difference, 5.7 tons, being just significant at the 5% level. The yields for variety 3 did not differ significantly from those of the other varieties at the two lower levels of N . At the highest level of N , the yield for variety 3 was significantly lower than that for either of the other varieties."

5.29 Interpretation of the Analysis: Second Example. This example is more complex, mainly because the factorial (3×3) is of an unusual type. The data come from a long-term experiment on meadow hay,

conducted at Lady Manner's School, Bakewell, England, with the cooperation of Rothamsted Experimental Station. The yields are for the 1937 season. The two factors are shown schematically below.

First factor (3 levels)		Second factor (3 levels)	
$\left\{ \begin{array}{l} \text{No manure} \\ \text{Mixed artificials} \\ \text{8 tons compost} \end{array} \right\}$	Applied in 1936	$\left\{ \begin{array}{l} \text{No manure} \\ \text{Mixed artificials} \\ \text{8 tons compost} \end{array} \right\}$	Applied in 1937

The nature of the experiment may become clear from a discussion of some of the individual treatment combinations. In any replicate there are nine plots, of which three received mixed artificial fertilizers in 1937. In the previous year, 1936, one of these three plots received no manure, one received mixed artificials, and one received compost. Thus these three plots enable us to compare the *residual* effects of the 1936 applications of artificials and compost, on plots which received artificials in the current year, 1937. Similarly, from other treatment combinations, we can compare the residual effects of artificials and compost on plots which were unmanured in the current year and on plots which received compost in the current year.

Further, in any replicate there are three plots which were unmanured in 1936. Of these, one had no manure, one had artificials, and one had compost in 1937. Consequently we may also assess the *direct* effects of artificials and compost applied in 1937 on plots which were unmanured in the previous year, and likewise on plots which received artificials or compost in the previous year. Both the artificials and compost contain the three common plant nutrients, nitrogen, phosphorus, and potash.

The system of treatments is an ingenious one, designed to measure direct and residual effects at the same time. The experiment was started in 1932, but for simplicity we will ignore any effects of treatments applied prior to 1936.

The treatment means and the analysis of variance are shown in tables 5.11 and 5.12. The analysis of variance was calculated from the treatment means, rather than from single plots, and the error m.s. has been adjusted so as to apply to a treatment mean. The direct effects are highly significant, the interactions are significant at the 5% level, but residual effects are not significant. In considering the nature of the effects leading to these results, it is again convenient to think in terms of simple effects rather than of main effects and interactions. It is worth noting that the standard error of a single entry in the two-way table is $\sqrt{6.656}$, or 2.58, while that for the difference between two entries is 3.65.

With regard to *residual* effects, table 5.11 shows that compost produced a large and highly significant increase, 13.4 cwt. \pm 3.65, on the plots that received no manure in the current year. On plots that received manure in 1937, either artificials or compost, there is no suggestion of any residual effect of compost. None of the residual effects of artificials approaches the level required for significance. These results are

TABLE 5.11 TREATMENT MEAN YIELDS (HAY, CWT. PER ACRE) (\pm 2.58)

Direct effects of 1937 treatments	Residual effects of 1936 treatments			
	None	Artificials	Compost	Mean
None	53.6	56.8	67.0	59.1
Artificials	80.8	82.3	80.5	81.2
Compost	74.3	69.1	70.0	71.1
Mean	69.6	69.4	72.5	70.5

TABLE 5.12 ANALYSIS OF VARIANCE OF TREATMENT MEANS

	d.f.	s.s.	m.s.
Direct effects	2	732.28	366.14 **
Residual effects	2	18.24	9.12
Interaction	4	97.01	24.25 *
Error	24	159.74	6.656

in accord with general fertilizer experience, since a compost is more likely to give residual effects than an inorganic fertilizer, and since residual effects would be expected to show up most clearly on plots which have no manures during the current year.

So far as *direct* effects are concerned, artificials were superior to compost whatever the residual treatment.

1936 manuring	1937 artificials - 1937 compost (cwt.)
No manure	+6.5
Artificials	+13.2
Compost	+10.5

The differences among these three figures are within the limits of experimental error. To test this point, we find the sum of squares of deviations of these differences, which comes to 22.73. This is divided by 2 to make it comparable with the analysis of variance, giving 11.36, which represents two of the 4 d.f. for interactions. The mean square, 5.68, is slightly below the error m.s. Thus the superiority of artificials appears to be consistent.

The increases for artificials and compost over no manure are much reduced on plots that received compost in the previous year.

1936 manuring	(1937 artificials — 1937 none)	(1937 compost — 1937 none)
Compost	+13.5	+3.0
Artificials and none (averaged)	+26.4	+16.5

The striking reduction on plots with 1936 compost is simply a reflection of the nature of the residual effect of compost as previously examined.

The conclusions from the experiment might be summarized as follows.

"Artificials applied in 1937 increased yields by 13.5 cwt. on plots that received compost in the previous year, and by 26.4 cwt. on plots that did not receive compost in the previous year. The corresponding increases due to 1937 compost were 3.0 cwt. and 16.5 cwt., respectively. The superiority of 1937 artificials over 1937 compost, which averaged 10.1 cwt., appeared to be independent of the type of manuring during the previous year.

"As regards residual effects, compost applied in 1936 increased yields by 13.4 cwt. on plots that were unmanured in 1937, but gave no apparent increase on plots manured in 1937. There were no significant residual effects of 1936 artificials."

It seems evident that the significant interaction m.s. in the analysis of variance must arise mainly from the fact that compost had a residual effect only when no manure was applied currently. As an exercise it may be instructive to isolate the part of the interaction s.s. that is due to this effect. Since artificials appeared to have no residual effect, we will combine the 1936 artificials with the 1936 unmanured plots. Consequently, the residual effects of compost are obtained from the comparison

$$2(1936 \text{ compost}) - (1936 \text{ artificials}) - (1936 \text{ none})$$

This quantity may be calculated separately for each of the 1937 treatments. Its values are

$$(1937 \text{ none}) = +23.6; \quad (1937 \text{ artificials}) = -2.1;$$

$$(1937 \text{ compost}) = -3.4$$

The contrast between the residual effect of compost on plots without 1937 manures and that on plots with 1937 manures may be estimated from the comparison

$$2(+23.6) - (-2.1) - (-3.4) = +52.7$$

The square of this quantity, with a suitable divisor, is a single component of the interaction s.s. Since the divisor for the quantities 23.6, etc., is 6, the divisor needed is 6×6 , or 36. Hence the sum of squares for this component is $(52.7)^2/36$, or 77.15. The remaining 3 d.f. for interactions have a sum of squares equal to 19.86. The mean square, 6.62, is no larger than the error m.s. This verifies the suggestion that the significance of the interaction m.s. can be attributed to the type of residual effect of compost.

Some readers may have difficulty in satisfying themselves that the component isolated above really is a part of the interaction s.s. It may be helpful to express the basic quantity, 52.7, as a linear function of the original mean yields in table 5.11. The multipliers of the means will be found to be as shown below.

1937 treatments	1936 treatments		
	None	Artificials	Compost
None	-2	-2	4
Artificials	1	1	-2
Compost	1	1	-2

The sums of the coefficients are zero over every row and column so that the expression is orthogonal to both sets of main effects. Hence it must be a part of the interaction. The coefficients also enable us to verify the divisor, 36, which is equal to the sum of the squares of the coefficients.

Since the interactions can be attributed to the behavior of a single group of plots (those with compost in 1936 and no manuring in 1937), it might be suspected that this treatment had been allotted by chance a favorable set of plots. The residual increase, 13.4 cwt., does seem rather large in relation to the direct effect of compost. An examination of previous results does not lend much weight to this suspicion. Over the 4 preceding years, the average residual response to compost, with no current manure, was 11.1 cwt. In 2 years, 1935 and 1933, the plots involved in this comparison were the same as those in 1937. For these years the average residual effect was 6.2 cwt. In the other 2 years, when a different set of compost plots is involved, the average residual effect was 16.0 cwt.

The two preceding examples are intended to illustrate the fact that the most informative subdivision of the treatment comparisons depends on the type of experiment. An analysis copied from a model that appears similar in form may be inappropriate or even meaningless. The experimenter should first decide which comparisons are necessary for the interpretation of the results. The subsequent analysis, either by *t*-tests

or by subdivision of the analysis of variance, should be directed towards these comparisons.

5.3 Designs for Factorial Experiments

5.31 Factorials in Complete Block Designs. Most types of experimental plan are suitable for factorial experiments. In particular, if the total number of treatment combinations is not large, the designs described in chapter 4 are frequently used. The relative advantages of complete randomization, randomized blocks, and latin squares are the same with factorial as with non-factorial sets of treatments. For illustration, we show arrangements for a 4×2 factorial (i) in 8 randomized blocks and (ii) in an 8×8 latin square. In the former case, only the first replicate is given.

TABLE 5.13 FACTORIAL EXPERIMENT ARRANGED IN (i) RANDOMIZED BLOCKS AND (ii) A LATIN SQUARE

8 replications		8 treatment combinations	
		<i>A</i> at 4 levels (1,2,3,4)	
		<i>B</i> at 2 levels (1,2)	
i. In randomized blocks		Analysis of variance	
Rep. I	Replications		7
12	Treatment combinations		7
31	<i>A</i> 3		
21	<i>B</i> 1		
32	<i>AB</i> 3		
42	Error		49
11	Total		63
22			
41			
ii. In a latin square		Analysis of variance	
42 11 22 12 31 41 32 21	Rows		7
21 32 31 41 22 12 11 42	Columns		7
12 41 42 11 32 21 22 31	Treatment combinations		7
31 22 21 42 11 32 41 12	<i>A</i> 3		
32 12 11 31 21 22 42 41	<i>B</i> 1		
41 42 32 22 12 31 21 11	<i>AB</i> 3		
11 21 12 32 41 42 31 22	Error		42
22 31 41 21 42 11 12 32	Total		63

In the next section, a worked example of the analysis of a large factorial experiment in randomized blocks is presented.

TABLE 5.14 WEIGHTS OF DENERVATED (y) AND CORRESPONDING NORMAL (x) MUSCLE (unit = 0.01 gram)

Rep. I		Number of treatment periods daily					
Length of treatment (minutes)	Type of current	One (a_1)		Three (a_3)		Six (a_6)	
		y	x	y	x	y	x
1 (b_1)	Galvanic	72	152	74	131	69	131
	Faradic	61	130	61	129	65	126
	60 cycle	62	141	65	112	70	111
	25 cycle	85	147	76	125	61	130
2 (b_2)	Galvanic	67	136	52	110	62	122
	Faradic	60	111	55	180	59	122
	60 cycle	64	126	65	190	64	98
	25 cycle	67	123	72	117	60	92
3 (b_3)	Galvanic	57	120	66	132	72	129
	Faradic	72	165	43	95	43	97
	60 cycle	63	112	66	130	72	180
	25 cycle	56	125	75	130	92	162
5 (b_5)	Galvanic	57	121	56	160	78	135
	Faradic	60	87	63	115	58	118
	60 cycle	61	93	79	126	68	160
	25 cycle	73	108	86	140	71	120
Rep. II							
1 (b_1)	Galvanic	46	97	74	131	58	81
	Faradic	60	126	64	124	52	102
	60 cycle	71	129	64	117	71	108
	25 cycle	53	108	65	108	66	108
2 (b_2)	Galvanic	44	83	58	117	54	97
	Faradic	57	104	55	112	51	100
	60 cycle	62	114	61	100	79	115
	25 cycle	60	105	78	112	82	102
3 (b_3)	Galvanic	53	101	50	103	61	115
	Faradic	56	120	57	110	56	105
	60 cycle	56	101	56	109	71	105
	25 cycle	56	97	58	87	69	107
5 (b_5)	Galvanic	46	107	55	108	64	115
	Faradic	56	109	55	104	57	103
	60 cycle	64	114	66	101	62	99
	25 cycle	59	102	58	98	88	135

5.32 Numerical Example: a $4 \times 4 \times 3$ Factorial in Randomized Blocks. A number of experiments have indicated that electrical stimulation may be helpful in preventing the wasting away of muscles that are denervated. A factorial experiment on rats was conducted by Solandt, DeLury, and Hunter (5.8) in order to learn something about the best type of current and the most effective method of treatment. The factors and their levels are shown below.

A	B	C
Number of treatment periods daily	Length of treatment (minutes)	Type of current
1	1	Galvanic
3	2	Faradic
6	3	60 cycle alternating
	5	25 cycle alternating

Treatments were started on the third day after denervation and continued for 11 consecutive days. There are 48 different combinations of methods of treatment, each of which was applied to a different rat. Two replications were conducted, using 96 rats in all.

The muscles denervated were the gastrocnemius-soleus group on one side of the animal, denervation being accomplished by the removal of a small part of the sciatic nerve. The measure used for judging the effects of the treatments was the weight of the denervated muscle at the end of the experiment. Since this depends on the size of the animal, the weight of the corresponding muscle on the other side of the body was included as a covariance variate. The data are shown in table 5.14.

For a covariance analysis we require analyses of variance for both the denervated muscle (y) and the normal muscle (x), and the analysis of the product (yx). From the discussion in section 5.27 and preceding sections, the reader should have little difficulty in completing the analysis. The replicate totals, treatment-combination totals, and the grand total are first computed. From the treatment totals, the three two-way tables are constructed, as given in table 5.15. Since each set of main effect totals is obtained twice, this part is self-checking (though only one set of main effect totals need be recorded). To illustrate the computations made on the $A \times B$ table, we have for the analysis of y

Total s.s. for $A \times B$ table

$$\begin{aligned}
 &= \frac{1}{8}[(510)^2 + (543)^2 + \cdots + (546)^2] - \frac{1}{96}[(6069)^2] = 1039 \\
 A &= \frac{1}{8 \cdot 2}[(1936)^2 + (2028)^2 + (2105)^2] - \frac{1}{96}[(6069)^2] = 447 \\
 B &= \frac{1}{2 \cdot 4}[(1565)^2 + (1488)^2 + (1476)^2 + (1540)^2] - \frac{1}{96}[(6069)^2] = 223
 \end{aligned}$$

By subtraction, the AB sum of squares is found to be 369. For the sum of products, we replace each y^2 by the corresponding yx product. Thus

$$\begin{aligned} \text{Total s.p.} &= \frac{1}{8}[(510)(1030) + (543)(977) + \cdots + (546)(985)] \\ &\quad - \frac{1}{96}(6069)(11,307) = 1151 \end{aligned}$$

Finally, the ABC sum of squares is found by calculating the total s.s. among all 48 treatment combinations, and subtracting the sum of squares for A , B , C , AB , AC , and BC .

TABLE 5.15 TWO-WAY TABLES IN A 4 × 4 × 3 FACTORIAL

		y				x			
		a_1	a_3	a_6	Total	a_1	a_3	a_6	Total
B Length of treatment	b_1	510	543	512	1,565	1,030	977	897	2,904
	b_2	481	496	511	1,488	902	1,038	848	2,788
	b_3	469	471	536	1,476	941	896	1,000	2,837
	b_5	476	518	546	1,540	841	952	985	2,778
C Type of current	G	442	485	518	1,445	917	992	925	2,834
	F	482	453	441	1,376	952	969	873	2,794
	60	503	522	557	1,582	930	985	976	2,891
	25	509	563	589	1,666	915	917	956	2,788
A totals		1,936	2,028	2,105	6,069	3,714	3,863	3,730	11,307

		y				x			
		b_1	b_2	b_3	b_5	b_1	b_2	b_3	b_5
C Type of current	G	393	337	359	356	723	665	700	746
	F	363	337	327	349	737	729	692	636
	60	403	395	384	400	718	743	737	693
	25	406	419	406	435	726	651	708	703

The analyses of y^2 , yx , and x^2 are given in table 5.16. The regression coefficient of weight of denervated on weight of normal muscle is $3977/16,013$, or 0.248361 . The next step is to form the analysis of $(y - bx)^2$. This is done by adding to any y^2 value $(-2b)$ times the corresponding yx value, plus (b^2) times the corresponding x^2 value. The mean squares for $(y - bx)$ are the quantities used to test the significance of treatment effects on the weights of denervated muscles, adjusted for their regression on the weights of the normal muscles. As shown in section 3.86, the treatment m.s. is slightly inflated because of the sampling error of the regression coefficient. The inflation is sufficiently small, however, that only effects which appear to be on the borderline of significance need be tested by the exact method (section 3.86), which is more laborious to calculate.

The means have been adjusted in the usual way for the regression on the weight of the normal muscle. Because of this adjustment, every mean has a slightly different standard error. However, as pointed out in section 3.85, it is sufficiently accurate to use an average standard error. This is computed from the *effective* error m.s., which is 48.1×1.019 , or 49.0, as described in section 3.85. Thus the standard error for an entry in the two-way table is $\sqrt{49.0/8}$, or 2.47.

The 25 cycle alternating current gave a significantly higher mean weight than any other type of current. The 60 cycle alternating current was superior to both galvanic and faradic currents, the last two not being significantly different. The weights increased as the number of treatment periods daily increased: On inspection the increase appears approximately linearly related to the number of treatment periods.

To test this supposition, we may fit a regression of the adjusted A totals on the number of treatment periods (the adjusted A means do not carry enough decimals to check with the analysis of variance). The relevant data are shown below.

	Adjusted A totals	z = number of periods	
a_1	1950	1	
a_3	2005	3	$\sum (a - \bar{a})(z - \bar{z}) = 421.667$
a_6	2115	6	$\sum (z - \bar{z})^2 = 12.667$

By the usual formula, the contribution of the regression to the sum of squares for A is $(421.667)^2/(12.667)(32)$, or 439. The last divisor, 32, is required because the A totals are totals over 32 rats. The sum of squares for regression accounts for practically all the sum of squares for A (441), so that there is no indication of deviation from linearity. The regression coefficient for a single rat is $421.667/(12.667)(32)$ or 1.0 unit. The conclusion is that, within the range of periods tested, each additional treatment period per day increased the weight by 1 unit.

The B means in table 5.17 do not indicate any consistent effect of length of an individual treatment period (1 to 5 minutes). The major part of the AC interaction seems to come from the anomalous behavior of the faradic current, which produced a drop in weight from a_1 to a_3 and a_6 . Since the interaction as a whole was not significant, it does not seem worth while to examine the statistical significance of this effect.

5.33 Other Designs. The total number of treatment combinations increases rapidly as the number of factors or the number of levels of a factor is increased. In this event the latin square necessitates an amount

of replication that is usually impracticable, while it becomes difficult to assemble homogeneous replications for a randomized blocks arrangement. Consequently, the experimental error per unit tends to increase. In order to avoid this increase in error, a number of designs have been produced. The basic principle, which is common to all these designs, is the use of a "block" that is smaller than a complete replication. That is, the arrangements are such that the differences among these "incomplete" blocks are eliminated from error, just as the differences among replicates are eliminated from error in a randomized blocks design. Unfortunately, as will be explained later, this reduction in block size can be accomplished only by the deliberate sacrifice of accuracy on certain treatment comparisons. The designs fall into three main groups according to the particular treatment comparisons that are sacrificed in this way.

In the *first group*, the treatment comparisons that are sacrificed are the high-order interactions. These designs are appropriate in lines of research where experience has shown that high-order interactions are nearly always negligible. The designs available in randomized incomplete blocks are described in chapter 6 and those which can be placed in latin squares in chapter 8.

In the *second group*, known in agriculture as split-plot experiments, a factor, or a group of factors and their interactions, are sacrificed. These designs, which are very widely used, are discussed in chapter 7. Finally, in certain cases it is possible to construct designs so that the loss in accuracy is spread evenly over all factors and their interactions. Within this group there are several types: balanced lattices (chapter 10), balanced incomplete blocks (chapter 11) and balanced lattice squares (chapter 12). In cases where the effects of the factors and the sizes of the interactions are rather unpredictable, these designs may be suitable.

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CHAPTER 6

CONFOUNDING

6.1 The Principle of Confounding

6.11 The 2^3 Factorial Experiment with Complete Confounding. This chapter deals with designs where the size of block is reduced by the sacrifice of accuracy on certain high-order interactions. The method will be illustrated first for a 2^3 experiment with three factors, A , B , and C , each at two levels. Since there are only 8 treatment combinations, the replication is not particularly large and in practice this experiment would most frequently be arranged in ordinary randomized blocks or perhaps in an 8×8 latin square. The example is chosen because of its simplicity.

The interaction of highest order is the ABC interaction. It will be recalled (section 5.23) that this interaction is estimated from the comparison

$$(abc) + (a) + (b) + (c) - (ab) - (ac) - (bc) - (1)$$

Suppose that each replicate in the experiment is divided into 2 blocks of 4 units each, such that one block contains abc , a , b , and c , while the other contains ab , ac , bc , and (1) . With 3 replicates the plan (before randomization) would be as follows.

TABLE 6.1 2^3 EXPERIMENT IN BLOCKS OF 4 UNITS, WITH ABC CONFOUNDED

Block	Rep. I 1	Rep. I 2	Rep. II 3	Rep. II 4	Rep. III 5	Rep. III 6
	abc a b c	ab ac bc (1)	abc a b c	ab ac bc (1)	abc a b c	ab ac bc (1)

There are two important properties of this plan. The total of blocks 1, 3, and 5, minus the total of blocks 2, 4, and 6, is the ABC interaction total. Thus the ABC interaction is one of the components of the com-

parisons amongst blocks. It is said to be *completely confounded* with blocks. Secondly, the other six factorial effects, A , B , C , AB , AC , and BC , will all be found to be orthogonal with the block totals. For instance, the AB interaction may be written (apart from the divisor)

$$[(abc) + (c) - (a) - (b)] + [(ab) + (1) - (ac) - (bc)]$$

Of the 4 units in any block, two carry a (+) sign in this expression and two carry a (−) sign. Consequently, if we increase all the observations in a selected block by any amount, say 50, the estimate of AB remains unchanged, and similarly for the other 5 factorial effects. This means that these 6 factorial effects are not influenced by differences amongst blocks. Various phrases are used to describe this property: the effects may be said to be unconfounded with blocks, or free from block effects, or to be composed entirely of *within-block* comparisons.

Thus differences amongst blocks of 4 units are eliminated from the experimental errors of the main effects and two-factor interactions, whereas with randomized blocks only differences amongst blocks of 8 units are eliminated. The reduction in effective block size is attained by making ABC the same as one of the block comparisons. There is no within-block information available about ABC .

The degrees of freedom in the analysis of variance separate as follows.

	d.f.
Blocks	5
A, B, C, AB, AC, BC	6
Error	12
	—
Total	23

All sums of squares are calculated in the usual way, so that there is no complexity in the computations.

The composition of the error s.s. is perhaps worth noting. Consider the 3 blocks (1, 3, and 5 in table 6.1) that contain the treatments abc , a , b , and c . These may be regarded as a randomized blocks experiment with 4 treatments and 3 blocks. Consequently the interaction of treatments with blocks contains six components. Similarly in blocks 2, 4, and 6 the interaction of the other set of treatments with blocks provides six components. These two sets of six components constitute the error for the complete experiment. In other words, the error term in a confounded factorial is made up of interactions between treatments and incomplete blocks. This remains true in the more complex designs that appear later in this chapter, even though in these cases the composition of the error is less easy to detect.

ABC does not appear explicitly in the analysis above. Actually, the experiment, if properly randomized, provides an estimate of error and a test of significance for ABC . In practice it is seldom worth while making this test, which is usually very insensitive. The test may, however, throw additional light on the nature of confounding. Let us ignore all other factorial effects and consider table 6.1 as the plan of an experiment to determine ABC , i.e., to determine the difference between the group of treatments abc, a, b, c and the group $ab, ac, bc, (1)$. From this point of view the block becomes the "experimental unit" and the experiment is of the ordinary randomized blocks type, having 2 "treatments" and 3 replicates. The total s.s. among the 6 "units" is, of course, the blocks s.s. in the previous analysis. Thus the blocks s.s. may be subdivided into:

	d.f.
Replicates	2
ABC	1
Error for ABC	2
	—
Total	5

If confounding has been effective, the error for ABC , being composed of comparisons among blocks, will be larger than the error (with 12 d.f.) which applies to the rest of the experiment. Moreover, the error for ABC is estimated from only 2 d.f., so that the test of ABC is a poor one. The test might be worth making with a considerable number of experiments of the same type, where it is desired to examine the average ABC effect over the whole group.

6.12 The 2^3 Factorial Experiment with Partial Confounding. Any of the seven factorial effects may be confounded with blocks in this way. The rest will then be free from block effects. These facts enable us to spread the confounding in an experiment among several factorial effects. A plan of this type is shown in table 6.2.

TABLE 6.2 2^3 EXPERIMENT IN BLOCKS OF 4 UNITS, WITH ABC, AC, BC PARTIALLY CONFOUNDED

	Rep. I		Rep. II		Rep. III	
Block	1	2	3	4	5	6
	abc a b c	ab ac bc (1)	abc ac b (1)	ab bc a c	abc bc a (1)	ab ac b c
	ABC		AC		BC	

The difference between the totals of blocks 1 and 2 represents the ABC interaction. In replication II, however, the block composition has been changed so that the difference between the block totals is the AC interaction. In replication III the difference is the BC interaction.

With this plan, A , B , C , and AB are entirely free from block effects. ABC is completely confounded with blocks in replicate I, but in the other two replicates the estimate of ABC is orthogonal with blocks. Thus a "within-block" estimate of ABC may be obtained from replicates II and III. Similarly, a "within-block" estimate of AC is available from replicates I and III, and one of BC from I and II. These three effects are *partially* confounded with blocks, since an estimate that is free from block effects can still be made for each effect. Each "within-block" estimate is derived from 2 out of the 3 replicates in the experiment. The ratio $\frac{2}{3}$ serves as a measure of the extent of the confounding. Yates (6.1) calls this ratio the *relative information* on the confounded interaction. The ratio gives the amount of information available on the partially confounded effect, relative to that available on an unconfounded effect.

In the analysis of variance, all factorial effects appear.

	d.f.
Blocks	5
A	1
B	1
C	1
AB	1
AC'	1'
BC'	1'
ABC'	1'
Error	11
Total	23

The sums of squares for blocks and for the unconfounded effects A , B , C , and AB are found in the usual way. The sum of squares for AC is calculated only from replicates I and III, the divisor for $[AC]^2$ being 16 instead of 24. Corresponding rules apply to BC and ABC . In the analysis of variance, the primes are inserted as a reminder that these effects are partially confounded and require special calculations.

6.13 Confounding in the 2^n Series. In order to use confounded designs intelligently, it is not necessary to understand in detail the methods by which they are constructed. The rules for the construction of confounded 2^n factorials will be briefly described, however, since they are

fairly simple and may help to explain the choice of plans presented at the end of this chapter.

If a replicate is to be divided into 2 blocks, *any* factorial effect can be confounded with blocks. In most cases we would confound the highest-order interaction in all replicates. Suppose that each incomplete block is to be further divided into 2, so that there will be 4 blocks (I ... IV) in a replicate. Having chosen one factorial effect to represent the comparison (I + II - III - IV), we may easily verify that *any other* factorial effect may be made to represent the comparison (I - II + III - IV). That is, to arrange a 2^n factorial in blocks of 2^{n-2} units, we may confound any two factorial effects that we choose.

The remaining block comparison (I - II - III + IV) represents a third factorial effect, which is also confounded with blocks. Barnard (6.2) has shown that this effect is uniquely determined by the two effects that were chosen. It is always their "generalized interaction," and is found by combining all the letters that appear in the two chosen factorial effects, and cancelling all letters that enter twice. Thus, if a 2^5 factorial is to be confounded in blocks of 2^3 , or 8, units, we might be inclined to choose *ABCDE* and *BCDE* as the first two factorial effects to be confounded. The generalized interaction of these two effects is *ABCCDDDEE* or, after cancellation, is *A*. Consequently, if these two effects are chosen for confounding, the main effect of *A* is also automatically confounded. These results may be summarized as follows. "If a replicate of a 2^n factorial is arranged in blocks of 2^{n-2} units, three factorial effects are confounded with blocks. Of these, two may be chosen at will: the third is their generalized interaction."

To proceed, a plan with 8 blocks in the replicate can be obtained from one with 4 blocks in the replicate by dividing each block into two parts. In making this division we may confound any factorial effect, except one of the three that is already confounded. Thus, in the 2^5 example, where *ABCDE*, *BCDE*, and *A* were confounded in the division into 4 blocks, we might choose say *ACE* for the division into 8 blocks. But with 8 blocks in the replicate, there are seven components of the comparisons among blocks, and each of these must represent one factorial effect. Hence, in addition to the four factorial effects that are known to be confounded, three others are confounded. These three are the generalized interactions of *ACE* with *ABCDE*, *BCDE* and *A*, or *BD*, *ABD*, and *CE*, respectively.

The general rule should be clear from this case. Suppose that a replicate in a 2^n factorial is to be divided into blocks of 2^{n-k} units, so that there will be 2^k blocks in the replicate. Then we may select any k factorial effects to be confounded, subject only to the restriction that none

of these must be a generalized interaction of any group of the others. A further $(2^k - k - 1)$ effects are automatically confounded. These are all the effects which can be expressed as generalized interactions of the group of effects selected for confounding.

In the construction of designs for practical use, it follows that the effects to be confounded must not be selected without examination of the other effects that are automatically confounded. If confounding is to be restricted to high-order interactions, we cannot choose $ABCDE$ and $BCDE$ in a 2^5 factorial, since A would then be confounded. In order to avoid confounding any main effects or first-order interactions in this case, a little trial will show that the only possible choice is two second-order interactions and one third-order interaction: for instance ABC , ADE , and their generalized interaction $BCDE$. The plans at the end of this chapter contain what appear to be the best choices for general use.

When the interactions to be confounded have been selected, there remains the problem of writing out the plan showing the treatment combinations that appear in each block. This may be done by the procedure used in the present discussion. The replicate is first divided into two by confounding one effect, then into four by confounding a second, and so on. The rules which express each factorial effect in terms of the original treatment combinations (section 5.24) decide the composition of the blocks at each stage. For experiments with a large number of factors, arranged in blocks of small size, Fisher (6.3) has given an alternative method that is more expeditious [see also Finney (6.4)]. In this method a group of 2^{n-k} letter combinations is constructed, such that the generalized interaction of this group with any treatment combination gives the members of the block that contains this treatment combination.

Fisher (6.3) has examined the general problem of confounding when the object is to keep all main effects and first-order interactions completely clear of block effects. He has shown that experiments up to the 2^7 factorial (128 treatment combinations) may be placed in blocks of 8 units, while blocks of 16 units can accommodate experiments up to the 2^{15} factorial (32,768 treatment combinations).

The preceding discussion applied to a single replication. With more than one replicate we may repeat the first replicate, in which case the confounding is complete for the interactions involved. Alternatively, the effects that are confounded may be changed from one replicate to the next, so that some intra-block information is available on all effects.

6.14 Example of a 2^4 Factorial Confounded in Blocks of 8 Units. Table 6.3 contains the plan and yields of a 2^4 field experiment on beans

TABLE 6.3 PLAN AND YIELDS (BEANS IN POUNDS) OF A 2^4 FACTORIAL EXPERIMENT

Rep. I					Rep. II				
Block <i>a</i> 375	<i>p</i>	<i>k</i>	<i>d</i>	<i>npk</i>	Block <i>a</i> 351	<i>npk</i>	<i>d</i>	<i>p</i>	<i>dnk</i>
	45	55	53	36		43	42	39	34
	<i>dnk</i>	<i>dnp</i>	<i>dpk</i>	<i>n</i>		<i>n</i>	<i>dnp</i>	<i>k</i>	<i>dpk</i>
	41	48	55	42		47	52	50	44
Block <i>b</i> 381	<i>dp</i>	<i>nk</i>	<i>dk</i>	<i>pk</i>	Block <i>b</i> 395	<i>nk</i>	<i>dp</i>	(1)	<i>np</i>
	50	44	43	51		43	52	57	39
	<i>dnpk</i>	(1)	<i>dn</i>	<i>np</i>		<i>pk</i>	<i>dk</i>	<i>dnpk</i>	<i>dn</i>
	44	58	41	50		56	52	54	42

conducted by the Rothamsted Experimental Station in 1936. The factors were

Dung (*D*): none, 10 tons per acre.

Nitrochalk (*N*): none, 0.4 cwt. N per acre.

Superphosphate (*P*): none, 0.6 cwt. P_2O_5 per acre.

Muriate of potash (*K*): none, 1.0 cwt. K_2O per acre.

With 16 treatment combinations in blocks of 8 plots, only one factorial effect is confounded in each replication. From the plan it will be seen that *DNPK*, the highest-order interaction, was confounded in each of the two replicates. The computations proceed as follows.

Step 1. Calculate the totals for each treatment combination, the block and replicate totals, and the grand total. The total s.s. and the sums of squares for replicates and for blocks within replicates should present no difficulty.

Step 2. This deals with the computation of the factorial effect totals. Write down each factorial effect total as a linear function of the treatment combination totals, as shown in table 6.4. The treatment combinations and their total yields are placed in a systematic order in the first two columns, and the remaining columns give the factorial effects. Although the writing of the table requires some time, the procedure is quite simple. When the columns for *D* and *N* have been written down, the column for *DN* is obtained as the *product* of corresponding signs in the *D* and *N* columns. Similarly, later in the table *DNP* may be obtained as the product of the signs of *D* and *NP*, or of *N* and *DP*, and so on. Thus it is necessary to know only the expressions for the main effects.

The factorial effect totals are shown at the bottom of each column.

TABLE 6.4 CALCULATION OF FACTORIAL EFFECT TOTALS IN A 2⁴ EXPERIMENT

Treatment combination	Total yield	Factorial effect																
		D	N	DN	P	DP	NP	DNP	K	DK	NK	DNK	PK	DPK	NPK	DNPK		
(1)	115	-	-	+	-	+	+	-	-	+	+	-	+	-	-	+	50	
d	95	+	-	-	-	-	+	+	-	-	-	+	+	+	+	-	-32	
n	89	-	+	-	-	+	-	-	-	+	-	+	+	-	-	+	-22	
dn	83	+	+	+	-	-	-	-	-	-	-	-	+	+	+	+	28	
p	84	-	-	+	+	-	-	+	-	+	+	+	-	-	-	-	18	
dp	102	+	-	-	+	+	+	-	-	-	-	-	-	+	+	+	-32	
np	89	-	+	-	+	-	+	-	-	+	-	+	-	-	-	+	-22	
dnp	100	+	+	+	+	+	+	+	-	-	-	-	-	+	+	+	28	
k	105	-	-	+	-	+	+	-	+	+	+	+	-	+	+	+	-32	
dk	95	+	-	-	-	-	+	+	+	-	-	-	-	-	-	-	18	
nk	87	-	+	-	-	+	-	-	-	+	+	+	-	+	-	+	-22	
dnk	75	+	+	+	-	-	-	-	+	+	+	+	-	-	-	+	28	
pk	107	-	-	+	+	-	-	+	+	+	+	+	+	-	-	+	-32	
dpk	99	+	-	-	+	+	+	-	+	-	-	-	+	+	+	+	-22	
npk	79	-	+	-	+	-	+	-	-	+	+	+	-	-	-	+	28	
dnpk	98	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	28	
	1502	-8	-102	32	14	88	50	8	-12	-14	-32	18	28	-22	-32	50		

The total for *DNPK* is not required, since owing to the confounding *DNPK* will not appear explicitly in the analysis of variance. The *DNPK* column, however, allows a check to be made, as follows. If the 15 totals are added, it may be verified by inspection of table 6.4 that their sum, +66, should equal 16 times the (*dnpk*) treatment total, minus the grand total, i.e.,

$$16(98) - 1502 = +66$$

Yates (6.1) gives a different method for obtaining the factorial effect totals. His method is completely automatic and does not require the expressions for the factorial effects to be written down. The present method, though probably not so speedy as Yates's method, is more flexible in that only those factorial effects that are of interest need be obtained. Sometimes the experimenter will wish to isolate only the main effects and two-factor interactions, previous experience having shown that interactions of higher order are likely to be negligible.

Step 3. The contribution of each factorial effect to the treatments s.s. in the analysis of variance is now obtained. Since there are 32 plots, the square of each effect is divided by 32. The analysis of variance appears in table 6.5. Note that *DNPK* is omitted, since it already appears

TABLE 6.5 ANALYSIS OF VARIANCE

	d.f.	s.s.	m.s.
Replications	1	3.1	
Blocks in reps.	2	123.2	61.6
<i>D</i>	1	2.0	
<i>N</i>	1	325.1**	
<i>P</i>	1	6.1	
<i>K</i>	1	4.5	
<i>DN</i>	1	32.0	
<i>DP</i>	1	242.0**	
<i>NP</i>	1	78.1	
<i>DK</i>	1	6.1	
<i>NK</i>	1	32.0	
<i>PK</i>	1	24.5	
<i>DNP</i>	1	2.0	
<i>DNK</i>	1	10.1	
<i>DPK</i>	1	15.1	
<i>NPK</i>	1	32.0	
Error	14	340.0	24.29
Total	31	1277.9	

in the blocks s.s. As a check, the total of the 15 treatment s.s., 811.6, plus that for *DNPK*, 78.1, should equal the treatments s.s. as found in

the ordinary way from the totals for the 16 treatment combinations. The error s.s. is found by subtraction.

Only two effects are significant: the main effect of nitrochalk, which produced a depression of the yield, and the *DP* interaction. Since neither *D* nor *P* produced significant main effects, it is advisable not to lay much stress on the *DP* interaction in the absence of other confirmatory evidence.

Step 4. The best presentation of the results will depend on their nature and on the audience for whom they are intended. Table 6.6 gives

TABLE 6.6 DIFFERENTIAL RESPONSES (POUNDS PER PLOT)

Factor	Mean response	Response with							
		Dung (<i>D</i>)		Nitrochalk (<i>N</i>)		Super. (<i>P</i>)		Mur. pot. (<i>K</i>)	
		Abs.	Pres.	Abs.	Pres.	Abs.	Pres.	Abs.	Pres.
Dung (<i>D</i>)	-0.5	-2.5	+1.5	-6.0	+5.0	+0.4	-1.4
Nitrochalk (<i>N</i>)	-6.4	-8.4	-4.4	-9.5	-3.2	-4.4	-8.4
Super. (<i>P</i>)	+0.9	-4.6	+6.4	-2.2	+4.0	-0.9	+2.6
Mur. pot. (<i>K</i>)	-0.8	+0.1	-1.6	+1.2	-2.8	-2.5	+1.0

s.e.: ± 2.46 for differential response; ± 1.74 for mean responses.

a concise presentation of the main effects and two-factor interactions in a form that has been used frequently at Rothamsted Experimental Station. The response to each factor is shown separately for each level of every other factor. Thus the row labelled "Dung" contains the mean response and the differential responses to dung. The figure -2.5 (response to dung with nitrochalk absent) is the average response to dung over all plots that did not receive nitrochalk. The table enables a quick appraisal to be made of the nature of the main effects and two-factor interactions. In the present case, of course, the table is scarcely needed because of the dearth of effects.

The data in the table are easily calculated from the factorial effect totals in table 6.4. For instance,

$$\text{Mean response to dung} = \frac{1}{16}[D] = -\frac{8}{16} = -0.5$$

$$\begin{aligned} \text{Mean response to dung} \\ \text{(nitrochalk absent)} &= \frac{1}{16}\{[D] - [DN]\} = -\frac{40}{16} = -2.5 \end{aligned}$$

$$\begin{aligned} \text{Mean response to dung} \\ \text{(nitrochalk present)} &= \frac{1}{16}\{[D] + [DN]\} = +\frac{24}{16} = +1.5 \end{aligned}$$

These relations may be verified from the signs in the columns in table 6.4.

The standard error per plot is $\sqrt{24.29}$, or 4.93. Since each differential response in table 6.6 is the difference between the means of two groups of 8 plots, its standard error is $(4.93)/2$, or 2.46. The standard error for a mean response is $\frac{1}{\sqrt{2}}$ times this value, or 1.74.

6.15 Confounding of a 3^2 Factorial. Let the symbol ij denote the treatment combination that has the i th level of A and the j th level of B ($i, j = 0, 1, 2$). As before, we wish to keep the main effects clear of block effects, that is, to confound only the AB interaction. Since there are 9 treatment combinations in a replicate and since 3 is the only factor of 9, the size of the incomplete block must be 3 units.

We have seen (section 5.26) that the main effects of A and B both contain two components, while the AB interaction has four components. Further, the main effects and the interaction can be divided into single components in various ways, and the division that is most appropriate for interpretation of the results will change from experiment to experiment. This suggests that the particular components of the interaction which we should desire to confound will vary from case to case. Actually, the possibilities for confounding are much more restricted with the 3^n than with the 2^n system: in fact, only one set of components of AB lends itself readily to confounding.

The confounding is based on the properties of the 3×3 graeco-latin square. Suppose that the 9 treatment combinations are set out as in table 6.7, on which a 3×3 square is superimposed. Following the notation

TABLE 6.7 USE OF A 3×3 GRAECO-LATIN SQUARE TO OBTAIN THE AB INTERACTION

	b_0	b_1	b_2
a_0	(00) I_1J_1	(01) I_2J_2	(02) I_3J_3
a_1	(10) I_3J_2	(11) I_1J_3	(12) I_2J_1
a_2	(20) I_2J_3	(21) I_3J_1	(22) I_1J_2

used by Yates (6.1), we denote the latin letters in the square by I_1, I_2 , and I_3 , respectively, and the greek letters by J_1, J_2 , and J_3 , respectively.

It is clear that comparisons among the row totals of the square give the two components of the main effect of A , while comparisons among the column totals give the main effect of B . Now consider the I totals:

$$I_1 = (00) + (11) + (22); \quad I_2 = (01) + (12) + (20);$$

$$I_3 = (02) + (10) + (21)$$

By the basic property of the latin square, comparisons among these totals are orthogonal to both rows and columns: that is, to the main effects of A and B . Hence, the comparisons among the I totals must represent two of the four components of the AB interaction. The same argument shows that the two remaining components of AB are derived from the comparisons among the J totals, where

$$J_1 = (00) + (12) + (21); \quad J_2 = (01) + (10) + (22);$$

$$J_3 = (02) + (11) + (20)$$

A numerical verification of this fact was made at the end of section 5.26, where the sum of squares for AB was calculated from the latin and greek letter totals which correspond to the I and J totals.

The application of this result to confounding is shown in table 6.8. There are two possibilities. In plan (i) the I components of AB are completely confounded with incomplete blocks, since the block totals have been made the same as the I totals. The main effects and the J components of AB are unconfounded. In plan (ii), the J components of AB are

TABLE 6.8 3×3 EXPERIMENT WITH AB PARTIALLY CONFOUNDED

Plan (i)			Plan (ii)		
Incomplete blocks			Incomplete blocks		
(i)	(ii)	(iii)	(i)	(ii)	(iii)
(00)	(01)	(02)	(00)	(01)	(02)
(11)	(12)	(10)	(12)	(10)	(11)
(22)	(20)	(21)	(21)	(22)	(20)
I_1	I_2	I_3	J_1	J_2	J_3
I components confounded			J components confounded		

completely confounded. It should be noted that neither the I nor the J components are easy to interpret in terms of the results of an actual experiment. They are selected for confounding because they are convenient for this purpose.

If the number of replicates in an experiment is even, we may use plan (i) in half the replicates and plan (ii) in the other half. With this arrangement the I components may be estimated (clear of block effects) from those replicates in which plan (ii) is used, and vice versa for the J

components. The relative accuracy is $\frac{1}{2}$ for each set of components. It may be shown that the relative accuracy is also $\frac{1}{2}$ for *any* component of AB in which we may be interested. The confounding is said to be *balanced* with respect to AB . Balanced confounding is preferable, for there is no reason to confound I more heavily than J .

This design has limited practical utility. It may be used where the factors are known to operate independently, so that AB will be negligible. If information about AB is wanted, a randomized blocks design is advisable unless there will be a substantial reduction in error variance from the use of incomplete blocks. The same principle of construction, however, leads to a very useful design for the 3^3 factorial.

6.16 Confounding of a 3^3 Factorial. The same notation will be used. Thus, (021) denotes the treatment combination $a_0b_2c_1$. Since there are 27 treatment combinations, the possible sizes of incomplete block are 3 and 9 units. The plan for blocks of 3 units necessitates confounding of the two-factor interactions and is not given here. With 9 units per block, only ABC need be confounded. In the case of the 3^2 factorial we were able to divide the 9 treatment combinations into groups of three (the I and J groups), such that comparisons among the group totals gave the components of AB . We shall show that the 27 treatment combinations in a 3^3 factorial can be divided into groups of *nine* such that comparisons among the group totals give the components of ABC . Each set of three groups will contribute two components of ABC . Since ABC has in all eight components or degrees of freedom, there will be four such sets.

The AB interaction can be calculated separately for each level of C . As before, we will obtain the AB interaction from its I and J components. Table 6.9 shows the I components for each level of C .

TABLE 6.9 THE I COMPONENTS OF AB SHOWN FOR EACH LEVEL OF C

	c_0	c_1	c_2
I_1	(000) + (110) + (220)	(001) + (111) + (221)	(002) + (112) + (222)
I_2	(010) + (120) + (200)	(011) + (121) + (201)	(012) + (122) + (202)
I_3	(020) + (100) + (210)	(021) + (101) + (211)	(022) + (102) + (212)

We may regard table 6.9 as a 3×3 table, in which each entry is the total of 3 treatment combinations. The row totals of the table give the I components of AB , while the column totals give the main effect of C . Further, we may take I and J totals from this table just as in table 6.7.

These totals will be called $I - I_1$, $I - J_1$, etc., since they come from the I components of AB . Thus

$$I - I_1: (000) + (110) + (220) + (011) + (121) + (201) + (022) + (102) + (212)$$

$$I - I_2: (001) + (111) + (221) + (012) + (122) + (202) + (020) + (100) + (210)$$

$$I - I_3: (002) + (112) + (222) + (010) + (120) + (200) + (021) + (101) + (211)$$

By the same argument as for the 3×3 design, the comparisons among these totals of 9 treatment combinations must represent two of the components of the interaction of AB with C ; that is, of the ABC interaction.

Consequently if we put all treatment combinations in $I - I_1$ into the first block, those in $I - I_2$ into the second, and those in $I - I_3$ into the third, we have a plan which completely confounds two of the eight components of ABC , and leaves all other factorial effects unconfounded. This arrangement appears as replication III in plan 6.7 for the 3^3 factorial at the end of this chapter.

The second set of three groups of 9 treatment combinations is obtained by taking the J totals from table 6.9.

$$I - J_1: (000) + (110) + (220) + (012) + (122) + (202) + (021) + (101) + (211)$$

$$I - J_2: (001) + (111) + (221) + (010) + (120) + (200) + (022) + (102) + (212)$$

$$I - J_3: (002) + (112) + (222) + (011) + (121) + (201) + (020) + (100) + (210)$$

This grouping constitutes replication IV in plan 6.7. The remaining sets are obtained by forming a 3×3 table similar to table 6.9 for the J components of AB at each level of C . The reader may verify that the $J - I_1$, $J - I_2$, and $J - I_3$ groups are as shown in replicate I and the $J - J_1$, $J - J_2$, and $J - J_3$ groups in replicate II of plan 6.7.

6.17 Example of a 3^3 Factorial Confounded in Blocks of 9 Units.

This experiment, conducted by the Seed Laboratory, Iowa State College, tests the effects of 3 levels of nitrogen, 3 of phosphorus, and 3 of potash on the germination of lettuce seedlings. The seed was thoroughly mixed, and divided into 108 samples of about 60 seeds each. Each sample was planted in a copper box, 6 in. square and $1\frac{1}{2}$ in. deep, in a mixture of soil and sand. The boxes were placed in a germinator, at a temperature of about 32°C . At the end of 5 to 7 days the seedlings were classified as normal, abnormal, hard, or dead. The data in table 6.10 show the numbers of normal lettuce plants.

There were 4 replications, each placed on a different shelf in the germinator. On a shelf the boxes were placed in 3 columns of 9 boxes each, each column being an incomplete block. It will be noted that all 3 fertilizers had a deleterious effect on emergence.

TABLE 6.10 NUMBER OF LETTUCE PLANTS EMERGING IN A 3³ EXPERIMENT

Blocks	Rep. 1			Rep. 2		
	A npt No.	B npt No.	C npt No.	A npt No.	B npt No.	C npt No.
	012 11	201 42	111 46	121 39	100 32	010 26
	122 11	121 20	102 48	-220 34	012 37	120 16
	-220 13	210 24	221 58	102 31	210 37	+200 6
	-202 12	011 38	001 53	201 34	221 33	021 13
	101 11	112 39	010 54	-000 40	-202 27	211 12
	021 30	100 61	212 37	-022 31	+020 21	+222 12
	-000 41	+002 40	-022 41	212 26	111 12	+002 16
	110 21	+222 46	120 25	011 35	001 30	112 19
	211 21	+020 44	+200 32	110 38	122 22	101 14
	171	354	394 919	308	251	134 693

Blocks	Rep. 3			Rep. 4		
	A npt No.	B npt No.	C npt No.	A npt No.	B npt No.	C npt No.
	101 26	+020 19	-220 19	112 37	+020 47	+222 15
	210 17	211 25	122 19	121 31	+002 33	201 42
	221 21	121 18	100 42	010 44	221 15	012 41
	-000 38	110 27	201 11	-022 25	122 29	210 24
	112 22	+222 20	010 36	100 39	011 38	-000 52
	011 19	001 40	111 37	211 28	212 27	111 28
	120 18	012 30	+002 29	-202 35	+200 38	120 30
	-202 22	102 28	021 27	001 37	101 31	102 50
	-022 14	+200 25	212 13	-220 26	110 32	021 29
	197	232	233 662	302	290	311 903
						Grand total 3177

The use of the + and - signs is discussed later in this section.

TABLE 6.11 ADDITIONAL SUMMARY TOTALS

Treatment totals				NPK components			
n_0			n_1	n_2			Total
	p_0	p_1	p_2	p_0	p_1	p_2	
k_0	171(000)	160(010)	131(020)	174(100)	118(110)	89(120)	101(200)
k_1	160(001)	130(011)	99(021)	82(101)	123(111)	108(121)	129(201)
k_2	118(002)	119(012)	111(022)	157(102)	117(112)	81(122)	96(202)
							92(220)
							127(221)
							93(222)
Two-way tables				NPK components			
	n_0	n_1	n_2	1A	1B	1C	Total
p_0	449	413	326	944	1102	1131	3177
p_1	409	358	291	171	354	394	919
p_2	341	278	312	773	748	737	2258
k_0	462	381	295	2A	2B	2C	Total
k_1	389	313	342	1119	1113	945	3177
k_2	348	355	292	308	251	134	693
	1199	1049	929	811	862	811	2484
				3A	3B	3C	Total
k_0	446	380	312	1025	1073	1079	3177
k_1	371	339	334	197	232	233	662
k_2	371	339	285	828	841	846	2515
	1188	1058	931	4A	4B	4C	Total
				1104	991	1082	3177
				302	290	311	903
				802	701	771	2274

The computations proceed as follows.

Step 1. Form the block and replicate totals and the grand total (shown in table 6.10) and also the totals for each treatment combination and the three two-way tables (shown in table 6.11).

Step 2. These data enable us to calculate the total s.s. and the sum of squares for replications, blocks within replications, and for the N , P , K , NP , NK , and PK factorial effects. All are obtained in the usual way and are entered in the preliminary analysis of variance (table 6.12).

TABLE 6.12 ANALYSIS OF VARIANCE FOR A $3 \times 3 \times 3$ FACTORIAL

	d.f.	s.s.	m.s.
Replications	3	2,041.88	
Blocks within replications	8	5,008.15	626.02
N	2	1,016.67	508.34 **
P	2	917.39	458.70 **
K	2	293.39	146.70
NP	4	399.27	99.82
NK	4	589.61	147.40
PK	4	212.89	53.22
NPK : confounded in replications			
1	2'	25.21	12.60
2	2'	64.22	32.11
3	2'	6.39	3.20
4	2'	198.30	99.15
Error	70	4,146.88	59.24
Total	107	14,920.25	

Subdivision of part of the treatments s.s.

N :	L	1	1,012.50	1,012.50 **
	Q	1	4.17	4.17
P :	L	1	917.35	917.35 **
	Q	1	0.04	0.04
K :	L	1	284.01	284.01 *
	Q	1	9.37	9.37
NP :	$L \times L$	1	184.08	184.08
	Rest	3	215.19	71.73
NK :	$L \times L$	1	256.69	256.69 *
	Rest	3	332.92	110.97
NPK :	$L \times L \times L$	1'	59.12	59.12

Notice that we do not calculate the total s.s. for treatments (with 26 d.f.), since this contains part of the blocks s.s.

Step 3. There remains the calculation of the contribution from the *NPK* interactions. Before doing this, it may be remarked that sometimes, from the nature of the factors or from previous experience, there is good reason to believe that the three-factor interactions will be negligible. In this case the experimenter may decide to pool the sum of squares for *NPK* with the error s.s. without troubling to compute the sum of squares for *NPK*. The pooled error would have 78 d.f. and would be obtained by subtraction of the sum of squares calculated in step 2 from the total s.s. The danger from this procedure is that if three-factor interactions are present the experimenter may not detect them, and moreover his estimate of error will be inflated. For many types of field experimentation in agriculture the procedure seems reasonably safe. When there is doubt it is better to isolate *NPK*.

Consider the two components of *NPK* that are confounded in replication 1. The contribution of these components to the sum of squares for *NPK* must be calculated from the remaining 3 replicates, in which they are unconfounded with blocks. The totals needed are shown in table 6.11 under the heading "*NPK* components." From the treatment totals, compute the total (944) of the 9 treatment combinations that appear in block 1A. Thus

$$\text{Block 1A} = 012 + 122 + 220 + 202 + 101 + 021 + 000 + 110 + 211$$

$$944 = 119 + 81 + 92 + 96 + 82 + 99 + 171 + 118 + 86$$

In the same way we obtain 1102 for the total over all treatments that appear in block 1B and 1131 for block 1C. Underneath these figures we place the respective totals for blocks 1A, 1B, and 1C. By subtraction we obtain the totals 773, 748, and 737. These are the totals of the groups of treatment combinations taken over replications 2, 3, and 4. The sum of squares of deviations of these quantities from their mean is divided by 27, since each figure contains 27 observations. The result, 25.21, is the contribution of the two components of *NPK* to the sum of squares for *NPK*. The remaining six components are found similarly from replications 2, 3, and 4. Of course, all eight components could be computed in one step as

$$\frac{1}{27}[(773)^2 + (748)^2 + \dots + (701)^2 + (771)^2] - \frac{1}{81}[(2258)^2 + \dots + (2274)^2]$$

The error s.s., with 70 d.f., is calculated by subtraction.

Step 4. In an experiment of this type, where each factor has equally spaced levels of an ingredient, it is usually advisable to examine the linear and quadratic components of the response curves. The contributions to the treatments s.s. are displayed in the lower section of table 6.12.

The method of calculation was described in section 5.26, where the data for N and P in this experiment were used as an example. All three fertilizers show significant linear responses, with no indication of any departure from linearity.

With regard to the two-factor interactions, the "linear by linear" components have been isolated for NP and NK (see section 5.26 for the method). For PK it was not thought worth while to do this, since the total s.s. for PK , 212.89, is not large enough to allow any single component to be significant. The "linear by linear" component is significant for NK but not for NP .

Where the "linear by linear" components of the two-factor interactions are large, it is desirable to isolate the "linear by linear by linear" component of the three-factor interaction. This will be done as an exercise. First we must define this component. With *two* factors, say N and P , the $N_L P_L$ component has been defined as

$$(22) + (00) - (20) - (02)$$

where the numbers refer to the levels of N and P . The value of this quantity at the highest level of K , *minus* its value at the lowest level of K , measures the linear effect of K on $N_L P_L$, and is the $N_L P_L K_L$ component in question. Algebraically, it is

$$L = (222) + (200) + (020) + (002) - (022) - (202) - (220) - (000)$$

The estimate from the treatment totals will be found to be -27 .

Since three-factor interactions are partially confounded with blocks, this component is also partially confounded and must be adjusted so as to remove block effects. The nature of the confounding is seen by writing down the expression above in the original table of plot yields (table 6.10). This has been done by means of the $+$ and $-$ signs that appear to the left of the seedling numbers in the table. It is evident that L is orthogonal to the totals of blocks 1C, 2B, 3C, and 4C, since each of these blocks has one $+$ and one $-$. In blocks 1B, 2C, 3B, and 4B, L has three $+$ signs, while in the remaining blocks it has three $-$ signs. Hence, a quantity that is free from block effects is

$$3L - (1B) - (2C) - (3B) - (4B) + (1A) + (2A) + (3A) + (4A)$$

or

$$3(-27) - 354 - 134 - 232 - 290 + 171 + 308 + 197 + 302 = -113$$

the block symbols denoting block totals. This expression, apart from a divisor, may be shown to be the least squares estimate of $N_L P_L K_L$, adjusted for block differences.

If this quantity is expressed algebraically as a linear function of the plot yields, the sum of squares of the coefficients is found to be 216. The

contribution of $N_L P_L K_L$ to the treatments s.s. is therefore $(113)^2/216$, or 59.12, which is practically the same as the error m.s. If the number of replicates differs from four, the procedure for isolating the "linear by linear by linear" component remains the same, except that the divisor for the square becomes $54r$ instead of 216.

Step 5. This concerns the presentation of the results. Usually it will be sufficient to show the three two-way tables of means, which are derivable from the two-way tables of totals (in table 6.11) on division by 12. Since all main effects and two-factor interactions are unconfounded, standard errors and t -tests for the 3×3 tables are obtained just as in a randomized blocks experiment. The principal results are that each fertilizer has produced a significant decrease in the numbers of seedlings that emerged, the decrease being substantially proportional to the amount of dressing. The significant $N_L K_L$ interaction represents the fact that the decrease in emergence from n_0 to n_2 was smaller at the k_2 level than at the k_0 level. There is some indication of a similar effect with N and P , though this did not attain significance.

The individual 27 treatment totals or means cannot be used as they stand for interpretative purposes, since they contain some block effects. A table of these totals or means will probably be unnecessary unless some aspect of the three-factor interaction requires study. To obtain such a table, we adjust each total so as to remove block effects. Each block effect is first estimated. For this purpose we do not use simply the observed block mean, since that in turn contains treatment effects. The least squares estimate of any block effect is

$$\frac{1}{27}[4(\text{block total}) - (\text{total of treatments appearing in the block})]$$

The basic data needed are available in the section headed "NPK components" in table 6.11. Thus, for block 1A, the estimated effect is

$$\frac{1}{27}[4(171) - (944)] = -9.6$$

The block effects are given below.

Replication	Block		
	A	B	C
1	-9.6	+11.6	+16.5
2	+4.2	-4.0	-15.1
3	-8.8	-5.4	-5.4
4	+3.9	+6.3	+6.0

To adjust any treatment total, we note from the plan the 4 blocks in which it appears, and compute the sum of the effects for these 4 blocks.

This quantity is *subtracted* from the unadjusted total to give the adjusted total. Thus for $n_1p_1k_0$, which appears in blocks 1A, 2A, 3B, and 4B, the adjusted total is

$$118 - [-9.6 + 4.2 - 5.4 + 6.3] = 122.5$$

To obtain the adjusted mean we divide by 4 as usual.

6.18 Mixed Series: the $3 \times 3 \times 2$ Factorial. In the designs discussed in previous sections, all factors have the same number of levels. When the number of levels differs from one factor to another, the utility of confounding is limited. Usually some two-factor interactions must be confounded, and the computations become more laborious. Only five designs of this type are given at the end of this chapter, though a number of others are available in the literature. As an example, we will consider the confounding of a $3 \times 3 \times 2$ factorial, which has 18 treatment combinations.

The main effects of a factor will be kept clear of block effects if every block contains each level of the factor the same number of times. Thus the factor *A*, which occurs at 3 levels (0, 1, 2), is unconfounded if every block contains an equal number of 0's, 1's, and 2's. Consequently, block size has to be a multiple of 3. Similarly we can keep the main effects of *B* (0, 1, 2) unconfounded if the block size is a multiple of 3. For the factor *C* at 2 levels (0, 1), the block size must be a multiple of 2. Hence, in order to keep all main effects clear of blocks, the only feasible block size is 6.

Further, with 6 units in a block, every possible combination of the levels of *A* and those of *C* can appear once in every block, so that we may expect to be able to keep *AC*, and likewise *BC*, unconfounded. We cannot place all the 9 combinations for *AB* in a block, so that *AB* will be partially confounded.

From this approach the plan can be constructed rather easily. Every block is to contain the 6 combinations (00), (10), (20), (01), (11), (21) of *B* and *C*. The only question is the manner in which the 3 levels of *A* are combined with the 6 pairs above. This allocation is to be such that all 6 combinations of *AC* appear in the block. That is, the 0, 1, and 2 levels of *A* must each appear with the 0 level of *C* and each with the 1 level of *C*. We may impose one additional rule, designed to confound *AB* as little as possible. Although we cannot represent all 9 combinations of *AB* in a block, it will be well to represent as many as possible. Thus we introduce the restriction that any *AB* combination (e.g., 12) must not appear more than once in a block.

Under these rules only four types of replicate can be made up, as shown in table 6.13.

Consider, for instance, in how many ways the block containing (000) can be constructed. The first three A levels (reading from the top in table 6.13) must be either 0, 1, 2 or 0, 2, 1, so that each A level shall appear with the 0 level of C . If 0, 1, 2 is chosen, the remaining three A

TABLE 6.13 POSSIBLE BLOCKS FOR A $3 \times 3 \times 2$ FACTORIAL

<i>B C</i>		Level of <i>A</i>											
		<i>Ia</i>	<i>Ib</i>	<i>Ic</i>	<i>IIa</i>	<i>IIb</i>	<i>IIc</i>	<i>IIIa</i>	<i>IIIb</i>	<i>IIIc</i>	<i>IVa</i>	<i>IVb</i>	<i>IVc</i>
0	0	1	2	0	2	0	1	1	2	0	2	0	1
1	0	2	0	1	0	1	2	0	1	2	1	2	0
2	0	0	1	2	1	2	0	2	0	1	0	1	2
0	1	2	0	1	1	2	0	2	0	1	1	2	0
1	1	0	1	2	2	0	1	1	2	0	0	1	2
2	1	1	2	0	0	1	2	0	1	2	2	0	1

levels must be either 1, 2, 0 or 2, 0, 1, these being the only sequences that make no AB combination appear twice. These sequences give blocks Ic and IIb respectively. Similarly, the choice of 0, 2, 1 for the first three A 's leads only to blocks IIIc and IVb.

In the same way, we find that there are only four possible blocks containing the treatment combination (001) and four containing (002.) The three sets of 4 blocks can be grouped into 4 separate replications (I, II, III, and IV in table 6.13).

As Yates (6.1) has shown, the plan can be written in a more condensed form (table 6.14) which exhibits the nature of the confounding.

TABLE 6.14 CONDENSED FORM OF THE PLAN FOR A $3 \times 3 \times 2$ CONFOUNDED FACTORIAL

Level of C	Blocks											
	I _a	I _b	I _c	II _a	II _b	II _c	III _a	III _b	III _c	IV _a	IV _b	IV _c
0	I_2	I_3	I_1	I_3	I_1	I_2	J_2	J_3	J_1	J_3	J_1	J_2
1	I_3	I_1	I_2	I_2	I_3	I_1	J_3	J_1	J_2	J_2	J_3	J_1

For example, in block Ia the 3 combinations that appear with the zero level of C are (01), (12), and (20). This is the group that forms the

I_2 component of AB (see section 6.15). It follows that in the first 2 replicates the I component of AB is partially confounded; the J component can be verified to be unconfounded. In replicates III and IV the J component is partially confounded, while the I component is clear of blocks. ABC is partially confounded in all replicates. The relative information on AB has been shown to be $\frac{7}{8}$ and that on ABC , $\frac{5}{8}$.

6.19 Numerical Example of the $3 \times 3 \times 2$ Design. Table 6.15 shows the plan and analysis of variance for an experiment on the response of young tung trees to fertilizers. The results of this experiment have been reported by Merrill, Kilby, and Greer (6.17). The general object was to

TABLE 6.15 DATA FOR $3 \times 3 \times 2$ FACTORIAL EXPERIMENT IN BLOCKS OF 6 UNITS

Ia	Ib	Ic		IIa	IIb	IIc	
100: 80	200: 78	000: 38		200:136	000: 43	100: 89	
210: 86	010: 55	110: 73		010: 56	110: 81	210: 87	
020: 70	120: 82	220: 75		120: 64	220: 90	020: 66	
201: 74	001: 67	101: 78		101: 95	201: 81	001: 91	
011: 82	111: 67	211: 51		211: 76	011: 61	111: 97	
121: 86	221: 57	021: 66		021: 71	121: 65	221: 60	
478	406	381	1,265	498	421	490	1,409
IIIa	IIIb	IIIc		IVa	IVb	IVc	
100: 86	200: 73	000: 66		200: 88	000: 53	100: 81	
010: 79	110: 76	210: 85		110:107	210: 66	010: 58	
220: 73	020: 97	120:101		020: 70	120: 92	220: 56	
201: 97	001:116	101:117		101: 79	201: 88	001: 90	
111: 79	211: 86	011:106		011: 92	111:109	211: 68	
021:113	121: 81	221:102		221: 96	021: 95	121: 67	
527	529	577	1,633	532	503	420	1,455

Analysis of variance

Source	d.f.	s.s.	m.s.
Replications	3	3,837	1,279
Blocks within replications	8	2,836	354
Levels of 8-8-6 in row (A)	2	1,116	558
Meals (B)	2	254	127
Levels of 8-8-6 side-dressed (C)	1	868	868 *
8-8-6 in row \times meals (AB)	4'	1,129	282
8-8-6 in row \times 8-8-6 side-dressed (AC)	2	2,995	1,498 **
Meals by 8-8-6 side-dressed (BC)	2	424	212
8-8-6 in row \times meals \times 8-8-6 side-dressed (ABC)	4'	1,016	254
Error	43	8,909	207
Total	71	23,384	

discover the type of fertilizer application that would stimulate the early growth so that the young trees would be ready for transplanting to commercial orchards by early autumn. The data shown are the mean heights in centimeters of the 12 tung trees on each plot. For convenience in following the computations, the plot yields follow the same order as in table 6.13. Note that factor *A* appears last in table 6.13 and first in table 6.15.

The three factors were as follows, all dressings being *per acre*.

A		B		C	
8-8-6 fertilizer applied in the row				8-8-6 fertilizer applied as side-dressing	
0	None	0	None	0	None
1	200 lb.	1	Tung meal (650 lb.)	1	200 lb.
2	400 lb.	2	Cottonseed meal (650 lb.)		

The side-dressing was placed in a shallow furrow on one side of the row about 8 inches from the tung seedlings. Factors *A* and *B* were applied at planting (March 13), factor *C* on June 3, and the height measurements were made on October 23. It should be noted that all three factors supply each of the common plant nutrients (*N*, *P*, and *K*).

In an experiment of this type, one might anticipate that two-factor interactions will be present. If, for example, 400 lb. of fertilizer applied in the row supply all the nutrients that the young tree can utilize, plots that receive this dressing will show no response to either factor *B* or *C*. On the other hand, plots that receive the zero level of *A* may respond to the dressings supplied in factors *B* and *C*. It would not be wise to confound any two-factor interactions heavily. Since, however, the soils available for experiments had been found to be very variable, it was decided to arrange the experiment in blocks of 6 plots. As we have seen, this will partially confound *AB*, though as the relative information is high ($\frac{7}{8}$), not much information is lost on this comparison even if the incomplete blocks prove ineffective.

Table 6.16 shows the computation sheet leading to the analysis of variance. The steps are as follows.

i. Calculate the block totals (denoted by *Ia*, *Ib*, etc., and shown in table 6.15), and the treatment totals (table 6.16). Calculate the total s.s. and the blocks s.s. The latter has been divided into two components—replications (3 d.f.) and blocks within replications (8 d.f.). This separation enables us to estimate the decrease in error variance due to the reduction in block size from 18 to 6 plots. The analysis of variance appears at the foot of table 6.15.

ii. Form the two three-way totals as shown. From the margins of the table for ($c_1 + c_0$), calculate the sum of squares for the main effects

TABLE 6.16 COMPUTATION SHEET FOR $3 \times 3 \times 2$ EXPERIMENT, BLOCKS OF 6 UNITS

(i) Treatment totals

	c_0			c_1		
	b_0	b_1	b_2	b_0	b_1	b_2
a_0	200	248	303	364	341	345
a_1	336	337	339	369	352	299
a_2	375	324	294	340	281	315

(ii) Two-way tables

	$c_1 + c_0$				$c_1 - c_0$			
	b_0	b_1	b_2	Totals	b_0	b_1	b_2	Totals
a_0	564(1)	589(4)	648(7)	1,801	+164(1)	+93(4)	+42(7)	299
a_1	705(2)	689(5)	638(8)	2,032	+ 33(2)	+15(5)	-40(8)	8
a_2	715(3)	605(6)	609(9)	1,929	- 35(3)	-43(6)	+21(9)	-57
Totals	1,984	1,883	1,895	5,762	162	65	23	250

(iii) From $(c_1 + c_0)$

$$\begin{aligned} I_1 &= (1) + (5) + (9) = 1,862 \\ I_2 &= (2) + (6) + (7) = 1,958 \\ I_3 &= (3) + (4) + (8) = 1,942 \end{aligned}$$

5,762

$$\begin{aligned} 2I_1' &= 2I_1 + I_a + II_a = 4,700 \\ 2I_2' &= 2I_2 + I_b + II_b = 4,743 \\ 2I_3' &= 2I_3 + I_c + II_c = 4,755 \end{aligned}$$

14,198

$$\begin{aligned} J_1 &= (1) + (6) + (8) = 1,807 \\ J_2 &= (2) + (4) + (9) = 1,903 \\ J_3 &= (3) + (5) + (7) = 2,052 \end{aligned}$$

5,762

$$\begin{aligned} 2J_1' &= 2J_1 + III_a + IV_a = 4,673 \\ 2J_2' &= 2J_2 + III_b + IV_b = 4,838 \\ 2J_3' &= 2J_3 + III_c + IV_c = 5,101 \end{aligned}$$

14,612

$$\text{s.s. for } AB = \frac{(4,700)^2 + (4,743)^2 + \dots + (5,101)^2}{84} - \frac{(14,198)^2 + (14,612)^2}{252} = 1,129$$

(iv) From $(c_1 - c_0)$

$$R_1 = +200, \quad R_2 = +32, \quad R_3 = +18; \quad S_1 = +81, \quad S_2 = +147, \quad S_3 = +22$$

where, e.g., $R_1 = (1) + (5) + (9), \dots, S_3 = (3) + (5) + (7)$

$$\begin{aligned} 2R_1' &= 2R_1 - I_b + I_c + II_b - II_c = +306 \\ 2R_2' &= 2R_2 + I_a - I_c - II_a + II_c = +153 \\ 2R_3' &= 2R_3 - I_a + I_b + II_a - II_b = +41 \end{aligned}$$

+500

$$\begin{aligned} 2S_1' &= 2S_1 - III_b + III_c + IV_b - IV_c = +293 \\ 2S_2' &= 2S_2 + III_a - III_c - IV_a + IV_c = +132 \\ 2S_3' &= 2S_3 - III_a + III_b + IV_a - IV_b = +75 \end{aligned}$$

+500

$$\text{s.s. for } ABC = \frac{(+306)^2 + (+153)^2 + \dots + (+75)^2}{60} - \frac{(+500)^2}{90} = 1,016$$

of A and B . From the *total* of the table for $(c_1 - c_0)$, we calculate the sum of squares for the main effect of C : this is $(250)^2/72$, since there are 72 plots. Similarly, the margins of the table provide the sum of squares for the AC and BC interactions, which being unconfounded are obtained in the usual way.

iii. These steps lead to the sum of squares for AB . First calculate the I and J components of AB , which are obtained by summation from the $(c_1 + c_0)$ table in step ii, as shown in table 6.16 (iii). Then apply the adjustments for block effects. Since I_1 , for instance, does not appear in blocks I_a or II_a , the adjustments involve the totals for these two blocks. The adjustments lead to the quantities $2I_1'$, etc., from which the sum of squares for AB follows as shown.

iv. These steps lead to the sum of squares for ABC . From the table for $(c_1 - c_0)$, step ii, form totals R_i and S_i which are calculated by the same rules as for the I 's and J 's, respectively. These totals are then adjusted for block effects, yielding quantities $2R_1'$, $2S_1'$, etc., from which the ABC s.s. is easily obtained. The error s.s. is then computed by subtraction.

v. Summary and presentation of results. Consider first the conclusions indicated by the analysis of variance. The only significant mean square for a main effect is that for C (side-dressing). The AC interaction is significant at the 1% level, while the mean square for A , though not significant, is substantially above the error m.s. Neither B (meals) nor any of its interactions approaches significance. These results suggest that the AC two-way table should be examined.

Since table 6.17 contains no confounded effects, the means per plot are found simply by addition and division from table 6.16, (i). It is

TABLE 6.17 AC TWO-WAY TABLE

Row dressing	Mean heights (centimeters) ± 4.15	
	Side dressing	
	c_0 None	c_1 200 lb.
a_0 None	62.6	87.5
a_1 200 lb.	84.3	85.0
a_2 400 lb.	82.8	78.0

evident that there are large and significant responses to the 200 lb. dressings, both when applied in the row and as side-dressing. The means, 84.3 for the row application and 87.5 for the side-dressing, do not differ significantly, so that the applications appear to have been about equally effective. There is no further increase to the 400 lb. row dressing, or to mixtures containing both types of dressing. Thus the large AC interaction arises primarily because each dressing is effective only in the absence of the other.

The analysis of variance would indicate that no definite effects of the meals (B) can be established. As an exercise, the AB two-way table, which involves partially confounded effects, will be examined. The means must be adjusted for block effects. This could be done by first calculating a mean for each incomplete block, adjusted for treatment effects, and then adjusting each AB mean for the 4 blocks in which it lies. It is slightly quicker to perform the adjustments by means of the I and J effects, as follows.

In table 6.16, (iii), divide the quantities $2I_1'$, etc., by 42, and calculate the deviations from their mean. The quantities $2J_1'$, etc., are treated similarly. The same procedure is carried out for the I 's and J 's, except that the divisor is 24. This gives

$$\begin{array}{llll} i_1' = -0.8, & i_2' = 0.2, & i_3' = 0.5 & j_1' = -4.7, \quad j_2' = -0.8, \quad j_3' = 5.4 \\ i_1 = -2.4, & i_2 = 1.6, & i_3 = 0.9 & j_1 = -4.7, \quad j_2 = -0.7, \quad j_3 = 5.5 \\ \delta i_1 = +1.6, & \delta i_2 = -1.4, & \delta i_3 = -0.4 & \delta j_1 = 0.0, \quad \delta j_2 = -0.1, \quad \delta j_3 = -0.1 \end{array}$$

The unadjusted mean of (a_0b_0) is 70.5. From table 6.16, (ii) and (iii), we see that (a_0b_0) belongs to I_1 and J_1 . Hence its adjusted value is

$$(a_0b_0)' = (a_0b_0) + \delta i_1 + \delta j_1 = 70.5 + 1.6 + 0.0 = 72.1$$

By the same principle, the adjustment to (a_2b_1) , for instance, is $(\delta i_2 + \delta j_1)$. The complete two-way table is shown below.

TABLE 6.18 AB TWO-WAY TABLE, ADJUSTED FOR BLOCK EFFECTS

Row dressing	Mean heights (centimeters)		
	b_0 None	b_1 Tung	b_2 Cottonseed
a_0 None	72.1	73.1	79.5
a_1 200 lb.	86.6	87.6	79.4
a_2 400 lb.	88.9	74.2	77.6

The standard error of a single figure in the table is $\sqrt{207/8}$, or 5.09, since each figure is the mean of 8 plots. This standard error applies to t -tests of unconfounded effects; i.e., of the main effects of A and B . For t -tests of components of the AB interaction, the figure must be multiplied by $\sqrt{8/7}$, or 1.069, the value $8/7$ being the inverse of the relative efficiency on AB .

For t -tests of quantities that are a mixture of main effects and interactions, the multiplier lies between 1.069 and 1. It will usually be sufficiently accurate to use the multiplier 1.069. If desired, the correct standard error may be obtained by means of a rule which will be given without proof. The quantity whose standard error is wanted is expressed in terms of components of main effects and interactions. Suppose that we wish to test the response to cottonseed when no row fertilizer is applied. The response is $(79.5 - 72.1)$, or 7.4 cm. The approximate standard error is $5.09 \times 1.069 \times \sqrt{2}$, or 7.69. To obtain the correct value, we write symbolically,

$$(b_2 - b_0)a_0 = \frac{(b_2 - b_0)(a_0 + a_1 + a_2)}{3} + \frac{(b_2 - b_0)(2a_0 - a_1 - a_2)}{3}$$

If all means are unadjusted means, this is an algebraic identity. The first term on the right is a component of the main effect of B , while the second is a component of the AB interaction. Further, by a property of least squares, the equation remains valid if means adjusted for block effects are substituted for unadjusted means. By the rule for the variance of a linear function, the variance of the first term on the right is $6\sigma^2/9$, where σ^2 is the variance of a single figure. By the same rule, the variance of the second term is

$$\left(\frac{12}{9}\right)\left(\frac{8}{7}\right)\sigma^2$$

the factor $8/7$ being introduced because the second term is a component of AB . The two terms are orthogonal, so that the estimated standard error of $(b_2 - b_0)a_0$ is

$$5.09\sqrt{\frac{6}{9} + \left(\frac{12}{9}\right)\left(\frac{8}{7}\right)} = 5.09\sqrt{2.190} = 7.53$$

Adjustment of the 18 individual treatment means is more tedious. In table 6.16, (iv), we divide the quantities $2R_1'$, etc., by 30, and the quantities R_1 , etc., by 24, thereafter taking deviations from the mean of each trio. This gives

$$\begin{array}{llllll} r_1' = +4.6, & r_2' = -0.5, & r_3' = -4.2 & s_1' = +4.2, & s_2' = -1.2, & s_3' = -3.1 \\ r_1 = +4.8, & r_2 = -2.2, & r_3 = -2.7 & s_1 = -0.1, & s_2 = +2.6, & s_3 = -2.6 \\ \delta r_1 = -0.2, & \delta r_2 = +1.7, & \delta r_3 = -1.5 & \delta s_1 = +4.3, & \delta s_2 = -3.8, & \delta s_3 = -0.5 \end{array}$$

In this experiment there is little point in adjusting all 18 treatment means. However, the unadjusted treatment totals, table 6.16, (i), suggest that there might be an effect of the meals (factor B) when applied alone. Hence it may be of interest to test the difference between the adjusted means of $a_0b_2c_0$ and $a_0b_0c_0$.

From table 6.16, (iii) and (iv), we see that $a_0b_0c_0$ belongs to i_1, j_1, r_1 , and s_1 . Since the unadjusted mean is 50.0, the adjusted mean is

$$50.0 + \delta i_1 + \delta j_1 - \delta r_1 - \delta s_1 =$$

$$50.0 + 1.6 + 0.0 + 0.2 - 4.3 = 47.5 \text{ cm.}$$

Note that the terms in δr and δs carry a *minus* sign whenever the treatment is at the c_0 level, and a *plus* sign when the treatment is at the c_1 level. For $a_0b_2c_0$, which belongs to i_2, j_3, r_2 , and s_3 , the adjusted mean will be found to be 73.1 cm.

In order to obtain the standard error of the difference ($73.1 - 47.5$) = 25.6, this must be expressed in terms of components of main effects and interactions. This requires some practice, but the method is not hard to master. Consider the following table of values of $(b_2 - b_0)$, averaged over the 4 replicates.

	$(b_2 - b_0)$			Total
	a_0	a_1	a_2	
c_0	d	e	f	l
c_1	g	h	k	m
Total	n	o	p	q

The quantity whose standard error we wish is d . Now it is easy to verify that

$$6d = q + (l - m) + (2n - o - p) + [(2d - e - f) - (2g - h - k)]$$

components of B BC AB ABC

As before, this is an identity for the unadjusted means and remains valid for adjusted means. Hence by the same rule as before

$$36\sigma_d^2 = \frac{\sigma^2}{4} \left[12 + 12 + 24 \left(\frac{8}{7} \right) + 24 \left(\frac{8}{5} \right) \right] = 22.46\sigma^2$$

where σ^2 is the estimated variance per plot. The extra factor $8/7$ is used for the AB component and the factor $8/5$ for the ABC component, for which the relative information is $5/8$. Consequently, the estimated standard error of d is

$$\sqrt{\frac{22.46 \times 207}{36}} = 11.4 \text{ cm.}$$

It follows that the t -value for the response to cottonseed when applied alone is $25.6/11.4$, or 2.25 , with 43 d.f., which is significant at the 5% level.

These results illustrate a feature that is common to the mixed series of confounded factorials. If the examination of the results is directed mostly at simple rather than factorial effects, the analysis is likely to be time-consuming because of the adjustments needed to remove block effects and to construct standard errors by the method given above. It will also be apparent that considerable experience in the analysis of variance, or access to expert advice on this topic, is advisable.

6.2 The Use of Confounded Designs

6.21 General Recommendations. In order to make efficient use of confounding, it is necessary to consider whether the advantage is likely to outweigh the disadvantages.

The advantage comes from the reduction in the experimental error by the use of a block which is more homogeneous, or which can be subjected to a more uniform technique, than the complete replicate. Without some idea of the amount of this reduction, a realistic decision cannot be made on the question of confounding. If uniformity data have been collected, the experimenter may compare the variability within the incomplete blocks with the variability within replicates. In addition, from the results of an experiment in which confounding has been employed, it is usually possible to estimate what the experimental error would have been if confounding had not been used. The method of calculation, which was first outlined by Yates (6.5, p. 215), is given in section 6.35. In many types of research the reduction in error from confounding varies greatly from experiment to experiment. In such cases little reliance can be placed on the results of a single trial; a summary of a moderately large group of experiments is necessary.

The disadvantages of confounding consist of (1) the reduction in replication on the confounded treatment comparisons, and (2) in some cases, a greater complexity in the calculations. With regard to the first, the experimenter should take into account the importance of the interactions that must be confounded and the extent to which they are confounded. No interaction should be *completely* confounded unless there is good reason to believe, either from previous experience or from the nature of the factors to be tested, that the interaction will be negligible.

As an illustration of the type of reasoning that might be employed in reaching a decision, consider the $3 \times 3 \times 2$ design discussed in the previous section. The incomplete blocks contain 6 units, with $7/8$ relative

information on AB and $5/8$ relative information on ABC . ABC is expected to be negligible; AB , however, is of sufficient interest so that confounding is undesirable if it involves any loss of precision for this comparison. The issue resolves itself into the question: is the error variance per unit with blocks of 6 units likely to be less than $7/8$ of the error variance with blocks of 18 units? If so, the AB interaction is estimated more precisely from the confounded design than from randomized blocks, because the reduction in variance per unit more than compensates for the loss of replication due to confounding. Since the relative information is useful for this type of argument, it is given in table 6.19 for all the plans presented at the end of this chapter.

The increase in the amount of computation varies both with the type of design and with the type of analysis needed to summarize the results. Where all the confounding is *complete*, the statistical analysis is practically the same as with randomized blocks. With all factors at two or three levels (i.e., in the 2^n and 3^n series), the extra computations are not formidable even when partial confounding is used; they are much more troublesome when the number of levels is not the same for all factors, as in the 3×2^2 and 3×2^3 designs. The extra work consists in the calculation of adjustments to the partially confounded comparisons so that their sums of squares in the analysis of variance are freed from block effects. If it is desired to present tables of mean yields which involve confounded comparisons, the entries in the tables must be adjusted so as to remove block effects. This point should be remembered, especially with designs where two-factor interactions are confounded. If t -tests of simple effects are needed, special computations must be made to obtain the standard errors for these tests. Moreover, if part of the data is missing, the analysis tends to become much more complicated than with randomized blocks.

The experimenter who has not previously used confounding and who does not have access to expert statistical advice is advised to confine himself at first to the simplest types of confounding. *He should make sure that he understands the method of analysis before the experiment is conducted.* After the experiment is completed, he should investigate the increase in precision that has been achieved by confounding, as a guide to the future use of the technique.

Yates (6.5) has pointed out that in certain circumstances a confounded design might give rise to a misleading interpretation of the results. This can be illustrated by means of the 2^3 factorial in table 6.1 (p. 154). This has three replicates, with ABC completely confounded in each replicate. Suppose that the response to A is much larger than any other effects, and that this response varies greatly from block to block. That is, the interaction of A with blocks is much larger than the interactions of the

other factors with blocks. Now the factorial effect total for BC may be written

$$BC = (abc) + (bc) + (a) + (1) - (ab) - (ac) - (b) - (c)$$

$$= [(abc) + (a) - (b) - (c)] - [(ab) + (ac) - (bc) - (1)]$$

$$= \text{response to } A \text{ in blocks 1, 3, 5} - \text{response to } A \text{ in blocks 2, 4, 6}$$

The point is that the factorial effect of BC is a component of the interaction of A with blocks. When tested against the error m.s. (12 d.f.) this response might be significant, not because there is an effect of BC , but because A has large interactions with blocks.

Another way of describing the source of the trouble is to say that the error m.s. is not homogeneous, in that the interaction of A with blocks is larger than the interactions of other treatments. If this were suspected, the danger of a misleading interpretation could be avoided by subdividing the error m.s. We can isolate 4 d.f. (two from blocks 1, 3, and 5 and two from blocks 2, 4, and 6) that represent the interaction of A with blocks. The mean square for these four could be used to test A and BC , while the remaining factorial effects would be tested by the other 8 d.f. for error. This device can usually be applied in the 2^n series, though generally not in the mixed series. Further, the heterogeneity of the error variance may not be suspected.

How frequently this danger arises will depend on the nature of the experimentation. Investigations by Yates (6.5) and Kempthorne (6.9) suggest that there is little danger with fertilizer experiments on English farm crops. In lines of research where errors have often been found to be heterogeneous, one should be on the alert for this effect.

In some experiments the use of confounding is practically unavoidable. For instance, an experiment on methods of manufacturing ice cream, planned at North Carolina State College, contained 3 factors each at 3 levels. It was desired to discover whether ordinary consumers could detect differences in the palatability of the 27 different types of ice cream. From previous experience it was believed that a taster (who was not an expert judge) could compare at most 3 ice creams at any single test. Moreover, it did not seem feasible to ask the tasters to grade the palatabilities on any kind of standard scale: they could merely be asked to rank the 3 ice creams in order of preference, if they had any preferences. Consequently, a design for a 3^3 factorial in blocks of 3 units was required, even though this involved confounding of two-factor interactions. The design adopted also employed partial confounding of the main effects, so as to obtain approximately equal accuracy on main effects and two-factor interactions.

TABLE 6.19 INDEX TO PLANS OF FACTORIAL EXPERIMENTS CONFOUNDED IN RANDOMIZED INCOMPLETE BLOCKS

a. Designs with which any number of replicates may be used

Type	Number of treatments	Number of units per block	Interactions confounded in a single replicate *	Plan
2^3	8	4	ABC	6.1
2^4	16	8	ABCD	6.2
2^4	16	4	AB, ACD, BCD	6.4
2^5	32	8	ABC, ADE, BCDE	6.5
2^6	64	16	ABCD, ABCE, CDEF	6.3
2^6	64	8	ABC, CDE, ADF, BEF, ABDE, BCDF, ACEF	6.6
3^3	27	9	ABC ($\frac{3}{4}$)	6.7
3^4	81	9	ABC, ABD, ACD, BCD, all ($\frac{3}{4}$)	6.8
4^2	16	4	AB ($\frac{2}{3}$)	6.12
4×2^2	16	8	ABC ($\frac{2}{3}$)	6.13

* The fractions in parentheses give the relative information on the comparisons which are confounded. Where no fraction is given, the comparison is completely confounded.

b. Balanced designs

Type	Number of treatments	Units per block	Number of replicates for a balanced design	Interactions confounded and relative information (in parentheses) *	Plan
2^4	16	4	$6n$ †	All two-factor ($\frac{5}{6}$); all three-factor ($\frac{1}{2}$)	6.4
2^5	32	8	$5n$	All three-factor ($\frac{4}{5}$); all four-factor ($\frac{1}{5}$)	6.5
2^6	64	8	$10n$	All three-factor ($\frac{4}{5}$); all four-factor ($\frac{1}{5}$)	6.6
3^3	27	9	$4n$	All three-factor ($\frac{3}{4}$)	6.7
3^4	81	9	$4n$	All three-factor ($\frac{3}{4}$)	6.8
3×2^2	12	6	$3n$ ‡	BC ($\frac{5}{6}$), ABC ($\frac{5}{6}$)	6.9
3×2^3	24	6	$3n$ ‡	BC, BD, CD, all ($\frac{5}{6}$); ABC, ABD, ACD, all ($\frac{5}{6}$)	6.10
$3^2 \times 2$	18	6	$4n$	AB ($\frac{1}{2}$), ABC ($\frac{5}{6}$)	6.11
4^2	16	4	$3n$	AB ($\frac{2}{3}$)	6.12
4×2^2	16	8	$3n$	ABC ($\frac{2}{3}$)	6.13
$4 \times 3 \times 2$	24	12	$9n$ ‡	AC ($\frac{2}{3}$), ABC ($\frac{2}{3}$)	6.14

* The factors ABC ... are read from the left; thus the BC interaction in a 3×2^2 design is the interaction between the 2 factors at 2 levels.

† The symbol " $6n$ " denotes that the number of replicates should be a multiple of 6.

‡ In these cases only the balanced design is recommended.

6.22 Index to Experimental Plans. The opportunities for confounding vary with the number of factors and with the number of levels of each factor. The plans given at the end of this chapter are those that are likely to be most frequently used.

Table 6.19, which forms an index to the plans, is divided into two parts. The first part shows a group of designs in which any number of replicates may be used. In successive replicates, the experimenter may either (i) repeat the first replicate (with a new randomization), in which case the same comparisons are confounded in all replicates, or (ii) change the grouping into blocks in successive replicates, so as to spread the confounding over a greater number of comparisons. In the plans, the factorial effects that are confounded are shown separately for each replication. Consequently, the reader can readily see what possibilities are open in this direction. Where confounding can be restricted to a high-order interaction, as in the 2^4 design in blocks of 8 units, the first procedure is preferable. On the other hand, with the 2^4 design in blocks of 4 units, where one two-factor interaction must be confounded in every replicate, the confounding should be changed in successive replications, unless it is certain that one of the two-factor interactions will be negligible. More detailed recommendations for the individual designs are given in section 6.3.

The second part, table 6.19b, shows a group of *balanced* designs, in which all interactions of the same order and the same type are confounded to the same extent. Thus in the $3 \times 2 \times 2 \times 2$ (or 3×2^3) design, all first-order interactions between the factors which have 2 levels are equally confounded, the relative information being 8/9. The first-order interactions between the factor at 3 levels and any factor at 2 levels are unconfounded. Balanced designs are useful in cases where all interactions of the same order and type are of equal interest, so that it is undesirable to confound some more fully than others. Also, the computations are usually simpler. For this reason, only the balanced designs should be used in certain cases, as indicated in table 6.19. On the other hand, with balanced designs the available numbers of replications are severely limited.

Table 6.20 shows a number of designs for which plans are given in the references cited. These designs are mainly for mixed series, and will not be discussed here.

6.23 Randomization. The procedure is the same for all designs.

1. Arrange the positions of the blocks at random within each replication, using a new randomization for each replication.

2. Allot the treatments in any block at random to the units in the block, using a new randomization for each block.

TABLE 6.20 CONFOUNDED DESIGNS FOR OTHER FACTORIAL EXPERIMENTS

Type	Units per block	Number of replicates	Interactions confounded	Reference
$3^2 \times 2^2$	12	$2n$	$AB (\frac{1}{8}), ABCD$	6.6
$3^3 \times 2$	18	$2n$	$ABC, ABCD$	6.6
$3^3 \times 2$	6	$2n$	$AB, AC, BC (\frac{1}{8}), ABD, ACD, BCD, ABC$	6.1
4×3^2	12	$2n$	$BC (\frac{1}{8}), ABC$	6.6
$4^2 \times 2$	16	Any	ABC	6.6
$4^2 \times 3$	12	$3n$	$AB (2\frac{1}{2}\frac{1}{7}), ABC$	6.6
4^3	16	Any	ABC	6.7
4×2^3	8	$3n$	ABC, ABD, ACD	6.6
4^4	16	Any	ABC, ABD, ACD, BCD	6.7
5^2	5	$4n$	$AB (\frac{3}{4})$	6.8
5×2^2	10	$5n$	$BC (2\frac{1}{2}\frac{1}{5}), ABC$	6.6
5^3	25	Any	ABC	6.8

6.24 Experiments in Single Replication. In most of the plans at the end of this chapter, the total number of treatment combinations is large. This fact limits the number of replications that can be employed. Sometimes the resources are sufficient for only a single replication unless the experimenter omits some factors which he originally intended to include.

With only one replication, we cannot derive an estimate of error from the interactions of treatments with blocks. If, however, certain high-order interactions are negligible, their mean squares in the analysis of variance will behave exactly like components of the error m.s., and therefore can be used to provide an estimate of error. In the 2^n series, the smallest design for which a single replicate is likely to be used is the 2^5 in blocks of 8 units (plan 6.5). In replicate 1 of this plan, ABC, ADE , and $BCDE$ are completely confounded. If the remaining three-, four-, and five-factor interactions furnish the estimate of error, the analysis of variance appears as follows.

	d.f.
Blocks	3
Main effects	5
Two-factor interactions	10
Error (from high-order interactions)	13
Total	31

Other designs in which a sufficient number of error d.f. are available from interactions among three or more factors are the 2^6 , the 3^3 , and the 3^4 . The interactions that constitute the estimate of error should be chosen before the results have been inspected. Recommendations for each of the designs above are given in the notes on the plans (section 6.3). It may sometimes be wise to deviate from these recommendations. For instance, in the 2^5 example above, the main effects of A , C , and D and the AC and AD interactions might all turn out to be rather large. This would suggest that the ACD interaction may not be negligible, and that it should not be included in the interactions used as error. It will be realized that a decision of this type should be made before examining ACD itself. The practice of examining the high-order interactions and using those that are small as error leads to a serious underestimation of the true error variance.

The estimation of error from high-order interactions is, of course, open to criticism. If some of these interactions happen to be large, the error m.s. that is used will overestimate the true error variance and the fact that the interactions are large will not be discovered. Where a considerable series of experiments of the same general type is being conducted, some safeguard is obtained both by examining the high-order interactions in experiments that are replicated and by watching the two-factor interactions in experiments with single replication. If most two-factor interactions are small, it seems very unlikely (though still possible) that interactions of higher order will be large. On the other hand, if many two-factor interactions are found to be large, this suggests that some three-factor interactions may also be substantial. For further discussion, see Yates (6.5) and Cornish (6.10).

6.25 Experiments with Fractional Replication. More recently several writers, Finney (6.11) and (6.12), Plackett and Burman (6.13), and Kempthorne (6.14) have considered the use of only part of a replication in a factorial experiment. Not enough experience has accumulated to permit appraisal of the utility of these designs, and only an introduction to them will be given. They are most likely to prove useful where (i) a large number of factors is tested, so that even a single replication is not feasible, and (ii) all high-order interactions are known with confidence to be negligible.

In order to see what happens when the experiment contains only part of a replication, consider a 2^3 factorial in which only the 4 treatment combinations a , b , c , and abc are tested. This is half of a complete replicate. It is clear that the three-factor interaction ABC cannot be estimated at all, since this represents the difference between the 4 treatments

that are tested and 4 that have not been tested. Finney (6.12) has called this comparison the *defining contrast*, since it defines the two halves into which the replicate is divided. Further, the main effect of A , as estimated from the difference between units with a and units without a ; i.e.,

$$(abc) + (a) - (b) - (c)$$

is exactly the same comparison as the BC interaction, as estimated from the difference between units that receive an even number of b 's or c 's and those that receive an odd number. Similarly, the estimate of B is the same as that of AC , while C is the same as AB . Factorial effects that are represented by the same treatment comparison are called *aliases*; thus BC is an alias of A . It has been shown (6.12) that the alias of any effect is its generalized interaction with the defining contrast. For example, the interaction of A with ABC is A^2BC or BC . This result makes it easy to discover the alias of any factorial effect.

If the experiment shows an apparent effect of A , there is no way of telling from the results whether the effect was due to A , or to the BC interaction, or to a mixture of the two. This type of information is too vague to be satisfactory. If, however, it can be taken for granted that all two-factor interactions are negligible, the ambiguity disappears, since we would conclude that an apparent effect of A was in fact due to A . With these assumptions, the half-replicate enables us to estimate the three main effects.

This example brings out the fact that in order to obtain definite information, it is necessary to assume that certain interactions are non-existent. The designs discussed by Finney and Kempthorne are adapted for situations where all interactions among three or more factors are negligible. In those presented by Plackett and Burman, even two-factor interactions are ignored.

We will examine the construction of a half-replicate of the 2^6 factorial, since this is probably the smallest member of the 2^n series for which fractional replication might be needed in practice. It is desired to obtain information on both main effects and two-factor interactions. First divide the treatment combinations into two sets of 32 by means of the defining contrast $ABCDEF$. Either set may be chosen as the half-replicate. The factorial effects are shown in table 6.21.

It will be noted that, rather fortunately, every alias of a main effect or two-factor interaction is a high-order interaction of a type that is assumed non-existent. The alias of a three-factor interaction is another three-factor interaction. Since all three-factor interactions are assumed negligible, these components may be used to estimate error. Consequently, this plan, if set out as a completely randomized design in 32

units, enables all main effects and two-factor interactions to be estimated and provides 10 d.f. from three-factor interactions for the estimation of error.

Since a completely randomized design is rather inaccurate for many types of experiment, an arrangement in blocks of 16 or 8 units may be preferable. If the 32 units are divided into two groups of 16, one fac-

TABLE 6.21 STRUCTURE OF A 2^6 FACTORIAL (HALF-REPLICATE)

Main effects	Alias	Main effects	Alias
A	BCDEF	D	ABCEF
B	ACDEF	E	ABCDF
C	ABDEF	F	ABCDE
Two-factor interactions	Alias	Two-factor interactions	Alias
AB	CDEF	BF	ACDE
AC	BDEF	CD	ABEF
AD	BCEF	CE	ABDF
AE	BCDF	CF	ABDE
AF	BCDE	DE	ABCF
BC	ADEF	DF	ABCE
BD	ACEF	EF	ABCD
BE	ACDF		
Three-factor interactions (used as error)	Alias	Three-factor interactions (used as error)	Alias
ABC	DEF	ACE	BDF
ABD	CEF	ACF	BDE
ABE	CDF	ADE	BCF
ABF	CDE	ADF	BCE
ACD	BEF	AEF	BCD

torial effect and its alias are confounded with groups. The best choice is one of the aliases from the set used as error, e.g., *ABC* and *DEF*. The plan is easily constructed. Suppose that the set of 32 which contains *abcdef* has been chosen as the half-replicate. This is divided into two groups of 16 such that one group (block 1) contains all treatment combinations carrying an *even* number of the letters *a*, *b*, or *c*. The plan and analysis are shown in table 6.22.

For a design in blocks of 8 units, we must confound with blocks another alias pair from the comparisons that are used as error. Suppose that *ABD* and its alias *CEF* are selected. It will be recalled (section 6.13) that the generalized interaction of this pair with *ABC* and *DEF* is also confounded. By the usual rule, this interaction is the alias pair *CD* and *ABEF*. It is unfortunate that a two-factor interaction is confounded, but further trial shows that this is unavoidable in blocks of 8

TABLE 6.22 A HALF-REPLICATE OF A 2^6 FACTORIAL IN BLOCKS OF 16 UNITS

Block 1	Block 2		d.f.
(1)	<i>ad</i>		
<i>de</i>	<i>ae</i>		
<i>df</i>	<i>af</i>		
<i>ef</i>	<i>bd</i>		
<i>ab</i>	<i>be</i>		
<i>ac</i>	<i>bf</i>	Blocks	1
<i>bc</i>	<i>cd</i>	Main effects	6
<i>abde</i>	<i>ce</i>	Two-factor interactions	15
<i>abdf</i>	<i>cf</i>	Error (from three-factorial interactions)	9
<i>abef</i>	<i>adef</i>		—
<i>acde</i>	<i>bdef</i>		
<i>acdf</i>	<i>cdef</i>	Total	31
<i>acef</i>	<i>abcd</i>		
<i>bced</i>	<i>abce</i>		
<i>bcd f</i>	<i>abcf</i>		
<i>bcef</i>	<i>abcdef</i>		

Note that the error d.f. are reduced from 10 to 9, since 1 d.f. has been assigned to blocks.

units. To construct the plan (table 6.23), divide each block of table 6.22 into halves that contain respectively an odd and even number of the letters *c* and *d*.

TABLE 6.23 A HALF-REPLICATE OF A 2^6 FACTORIAL IN BLOCKS OF 8 UNITS

Block					d.f.
1	2	3	4		
(1)	<i>de</i>	<i>ae</i>	<i>ad</i>	Blocks	3
<i>ef</i>	<i>df</i>	<i>af</i>	<i>bd</i>	Main effects	6
<i>ab</i>	<i>ac</i>	<i>be</i>	<i>ce</i>	Two-factor interactions	14
<i>abef</i>	<i>bc</i>	<i>bf</i>	<i>cf</i>	Error	8
<i>acde</i>	<i>abde</i>	<i>cd</i>	<i>abce</i>		—
<i>acdf</i>	<i>abdf</i>	<i>abcd</i>	<i>abcf</i>		
<i>bced</i>	<i>acef</i>	<i>cdef</i>	<i>adef</i>	Total	31
<i>bcd f</i>	<i>bcef</i>	<i>abcdef</i>	<i>bdef</i>		

In practice, the letters should be allocated to the factors so that *CD* is the two-factor interaction that is considered least important. In the analysis of variance, the sum of squares for blocks, main effects, and the unconfounded two-factor interactions are all computed in the usual way.

Other experiments for which plans are described in reference (6.12) are a half-replicate of a 2^7 factorial in blocks of 8 units, a half-replicate of a 2^8 factorial in blocks of 16, a quarter-replicate of a 2^8 factorial in blocks of 16, and a third-replicate of a 3^5 in blocks of 9.

For the case where two-factor interactions are ignored, Plackett and Burman (6.13) give methods for constructing a large number of plans in which all main effects can be estimated. The only restriction is that the number of units tested is a multiple of 4. For instance, there is a plan for a 2^7 factorial, using only 8 units (one-sixteenth of a replicate). All seven main effects can be obtained, though no comparisons remain for an estimate of error. There are further plans in 12, 16, 20, etc., units, giving 4, 8, 12, etc., d.f. for error. All these plans are set out for a completely randomized design: with further confounding a reduced block size could be employed. As an illustration the plan with 16 units is given below. To facilitate study of this plan, the treatment combinations are shown both in our usual notation and by means of + and - signs to denote the two levels of each factor.

Every factor occurs 8 times at each level. For every pair of factors, each of the sign combinations, ++, +-, -+, and --, occurs four times. Thus the estimates of all seven main effects are mutually orthogonal, and are calculated in the usual way. With complete randomization, 8 d.f. are available for the estimation of error.

TABLE 6.24 2^7 FACTORIAL IN 16 UNITS (ONE-EIGHTH REPLICATE)

Unit	Our notation	Treatment combination + and - notation						
		<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
1	(1)	-	-	-	-	-	-	-
2	<i>dg</i>	-	-	-	+	-	-	+
3	<i>efg</i>	-	-	+	-	-	+	+
4	<i>cdf</i>	-	-	+	+	-	+	-
5	<i>bef</i>	-	+	-	-	+	+	-
6	<i>bdefg</i>	-	+	-	+	+	+	+
7	<i>bceg</i>	-	+	+	-	+	-	+
8	<i>bcde</i>	-	+	+	+	+	-	-
9	<i>ae</i>	+	-	-	-	+	-	-
10	<i>adeg</i>	+	-	-	+	+	-	+
11	<i>acefg</i>	+	-	+	-	+	+	+
12	<i>acdef</i>	+	-	+	+	+	+	-
13	<i>abf</i>	+	+	-	-	-	+	-
14	<i>abdfg</i>	+	+	-	+	-	+	+
15	<i>abce</i>	+	+	+	-	-	-	+
16	<i>abcd</i>	+	+	+	+	-	-	-

Since it is not clear how frequently all factors can be assumed to be additive in their effects, the utility of these designs remains to be seen. As Yates (6.5) has pointed out, one case where there is additivity occurs when each factor represents an object to be weighed. Suppose that weighings are done on a chemical balance. Testing the $+$ level of a factor corresponds to placing the object in the right-hand pan; testing the $-$ level corresponds to placing the object in the left-hand pan.

Thus the rows of table 6.24 provide a set of instructions for performing 16 successive weighings on 7 objects. In the first weighing, all objects are placed in the left-hand pan; consequently the weight that is read is the total weight of all 7 objects. In the second weighing objects d and g are placed in the right-hand pan and the others in the left-hand pan, and so on. This series of 16 weighings enables us to estimate the weight of each object as precisely as if all weighings had been devoted to that object alone. To make the analogy complete, a balance with a zero error would be needed, the zero error corresponding to the mean yield of an experiment, which can also be estimated from the 16 results. For a discussion and references, see Mood (6.15).

6.3 Notes on the Plans and Statistical Analysis

6.31 2^n Series. In plans 6.1 to 6.6, one level of a factor is denoted by the corresponding letter; the other level by the absence of the letter. With designs that are not balanced, care must be taken to allocate the letters a, b, c, \dots to the factors in the experiment so that the interactions which are confounded are those that the experimenter has decided to confound. For instance, the first replication of plan 6.4 (2^4 design in blocks of 4) confounds the interaction AB , but no other two-factor interaction. The factors labelled A and B should be those whose interaction is of least interest.

The statistical analysis follows the same general method for all 2^n designs. The total s.s. and the sum of squares among blocks are obtained by the usual methods. The factorial effects may be computed by one of the methods outlined in section 6.14. When this calculation has been completed, the sums of squares are added for all unconfounded comparisons. Let this total be S_T . If all the confounding is complete, S_T is the treatment s.s., since the effects that are confounded appear already in the block s.s. For each comparison which is partially confounded, the sum of squares is calculated separately *from those replications in which the comparison is unconfounded*. The replications to be used are clear from the plans, which show the factorial effects that are confounded in each

replication. The total of these sums of squares is added to S_T to give the treatment s.s. The error s.s. is found by subtraction.

2³ factorial, blocks of 4 units. Plan 6.1, which confounds ABC completely, may be used for every replication. For a worked example, see reference (6.1), p. 19.

Alternatively, with 4, or a multiple of 4, replicates, we may confound partially all two-factor interactions and the three-factor interaction, the relative information on each comparison being $3/4$. The plan and statistical analysis are described in reference (6.1), p. 21.

2⁴ factorial, blocks of 8 units. Plan 6.2 should be used for every replication, $ABCD$ being completely confounded.

Note that the treatment s.s. contains only 14 d.f. since the single degree of freedom for $ABCD$ is included in the blocks.

2⁴ factorial, blocks of 4 units. Plan 6.4 gives a balanced design with 6 replications. If a smaller number of replicates is used, the replicates chosen depend on the interactions which the experimenter decides to confound. Since one two-factor interaction is confounded in each replicate, it will usually be preferable to confound different comparisons in successive replications. For instance, if AB and CD are to be confounded, replicates 1 and 6 should be selected. This arrangement gives $1/2$ relative information on AB , CD and the three-factor interactions ACD , BCD , ABC , ABD , all other comparisons being clear of blocks.

With a 2^4 experiment, the investigator may be uncertain whether to confound in blocks of 8 units or in blocks of 4 units. The former design is recommended unless (a) the interactions which are confounded with blocks of 4 units are known to be small or (b) the block of 4 units is much more homogeneous than the block of 8 units. The latter condition may apply where the experimental material is extremely variable, or falls naturally into groups of 4 units.

2⁵ factorial, blocks of 8 units. The balanced design (plan 6.5) requires 5 replicates. When fewer than 5 replicates are used, it may be satisfactory to confound completely two three-factor interactions and one four-factor interaction. In this case replication 1 is repeated the required number of times.

If some information is desired on all treatment comparisons, replications 1, 2, ... may be used up to the proposed number. With replications 1 and 2, for instance, the interactions ABC , ADE , ABD , BCE , $BCDE$, and $ACDE$ are partially confounded, each with relative information $1/2$.

The statistical analysis is described in reference (6.1), p. 27, for the case in which the experiment contains only a single replication.

2⁶ factorial, blocks of 16 units. Plan 6.3, which shows a single replication with $ABCD$, $ABEF$, and $CDEF$ confounded, may be used for all

replications. The 3 interactions above are completely confounded, and are omitted from the treatment s.s. in the analysis of variance.

If there is only 1 replication, an estimate of error may be obtained from the unconfounded four-, five-, and six-factor interactions. These comparisons supply 22 d.f., of which three are confounded with blocks, leaving 19 for the estimate of error. The separation of the 63 d.f. is then as follows:

	d.f.
Blocks	3
Main effects	6
Two-factor interactions	15
Three-factor interactions	20
Error (from high-order interactions)	19

2⁶ factorial, blocks of 8 units. In plan 6.6, the size of block is reduced from 64 to 8 units without confounding any two-factor interactions. Four of the 20 three-factor interactions and 3 of the 15 four-factor interactions are confounded in each replication. The balanced design requires 10 replications, i.e., 640 experimental units, and necessitates a large experiment.

With a single replication, the 63 d.f. in the analysis of variance subdivide as follows:

	d.f.
Blocks	7
Main effects	6
Two-factor interactions	15
Three-factor interactions	16
Error (from high-order interactions)	19

The simplest procedure when there is more than one replication is to confound completely a single set of 7 factorial effects by the repeated use of replication 1. For partial confounding, the successive replicates in plan 6.6 may be taken. In the first 4 replicates no three-factor interaction is confounded more than once.

6.32 3ⁿ Series. From plan 6.7 onwards, the levels or variations of each factor are denoted by the figures 0, 1, 2, ..., while the factors are read from left to right. The treatment 102 implies the middle level of the first factor, the lowest level of the second factor, and the highest level of the third factor. In an experiment the numbers 0, 1, 2, ... may be assigned to the levels or variations of a factor according to any convenient scheme.

The testing of three instead of two variations of each factor makes possible a more thorough evaluation of the effects of the factors. In

particular, the 3^3 factorial experiment has been much used in investigations of the responses to fertilizers, since the effects of each of 3 different plant nutrients may be explored at 3 levels of application, which provide some information on the shape of the response-curve.

3³ factorial, blocks of 9 units. In plan 6.7, two of the 8 d.f. for ABC are completely confounded in each replication, so that the balanced design requires 4 replicates and provides $3/4$ relative information on ABC . Unless the three-factor interactions are known to be negligible, in which case replication 1 may be used for all replicates of the experiment, it is preferable to select the required number of replications in succession from the plan. For example, an experiment with 3 replicates gives $2/3$ relative information on 6 of the 8 interaction d.f., and full replication on the remaining 2 d.f. As usual, the rule in the analysis of variance is to estimate confounded comparisons from those replications in which they are not confounded. An example of the analysis with 4 replicates has already been given (section 6.17).

For an example of the analysis with a single replicate, see reference (6.1), p. 53. If the three-factor interactions are used as error, the separation of the degrees of freedom is as shown on the left in table 6.25.

TABLE 6.25 SUBDIVISION OF THE DEGREES OF FREEDOM IN A 3^3 EXPERIMENT WITH SINGLE REPLICATION

I. Three-factor interactions used as error		II. For experiments where all effects are approximately linear	
	d.f.		d.f.
Blocks	2	Blocks	2
Main effects	6	Main effects	6
Two-factor interactions	12	AB : linear \times linear	1
Error	6	AC : linear \times linear	1
	—	BC : linear \times linear	1
Total	26	Error	15
		Total	26

The number of error d.f., 6, is rather small. There is an alternative scheme for experiments where the effects of the factors are linear, or nearly so. Under these circumstances, the only components of the two-factor interactions that are likely to be substantial are the interactions of the linear responses to the factors. Hence, nine of the 12 d.f. for two-factor interactions are added to the material used for an estimate of error, as shown in table 6.25, II. If in addition the linear components of the two-factor interaction are large, a further variant is to remove from the error the ABC : linear \times linear \times linear term. This leaves 14 d.f.

for error. The calculation of these components should be fairly simple with the help of the example worked in section 6.17.

3⁴ factorial, blocks of 9 units. With 4 factors each at 3 levels (plan 6.8) the block size is reduced from 81 to 9 units by confounding three-factor interactions. There are 4 three-factor interactions: *ABC*, *ABD*, *ACD*, and *BCD*, each with 8 d.f. Two components from each set of 8 are confounded in any replication. If the experiment has more than 1 replicate, partial confounding is generally preferable, the required number of replicates being selected from the balanced set. The balanced design, with 4 replicates, gives threefold replication on all three-factor interactions.

Statistical analysis with a single replication. An analysis may be calculated without difficulty from the following partition of the degrees of freedom.

	d.f.
Blocks	8
Main effects	8
Two-factor interactions	24
Error (from three- and four-factor interactions)	40

All sums of squares are calculated in the usual way. For the main effects and two-factor interactions, six 3×3 tables are required. The error, which is found by subtraction, contains 24 d.f. representing unconfounded three-factor interactions and the 16 d.f. for the four-factor interactions.

This analysis gives no test of the three-factor interactions. If a test of *ABC*, for example, is wanted, the first step is to calculate in the usual way the sum of squares for the 8 d.f. From this we must subtract the sum of squares for the 2 d.f. that are confounded with blocks.

In plan 6.8, replication I, the 2 d.f. are denoted by *ABC IV*, where the numeral IV refers to replication IV of plan 6.7. If we ignore *D*, it will be found that blocks 1, 6, and 8 contain the same set of 9 treatments as block IVa in plan 6.7. Calculate the total yield of these 3 blocks (27 units). Similarly, we calculate the total yield of blocks 2, 4, and 9, which correspond to block IVb in plan 6.7, and of blocks 3, 5, and 7, which correspond to block IVc. The sum of squares of deviations of the 3 totals, divided by 27, is subtracted from the sum of squares for the 8 d.f. The resulting 6 d.f. are removed from the error, which now has 34 d.f.

The reader may verify that the 2 confounded degrees of freedom for *BCD* (i.e., *BCD III*) are obtained from the totals of blocks 1, 2, 3, blocks 4, 5, 6, and blocks 7, 8, 9. In this case *A* is ignored when plan 6.8 is compared with plan 6.7.

Statistical analysis with more than one replication. If all three-factor interactions are presumed to be negligible, they may be combined with

the error. The sums of squares for blocks, main effects, and two-factor interactions are found in the usual way while the remainder constitutes the error.

All 32 components of the three-factor interactions may, however, be calculated and tested. Partially confounded comparisons are taken only from replicates in which they are unconfounded. For example, if there are two replications of plan 6.8, all the data are used in the calculation of components *ABC* I and *ABC* II, which are unconfounded. For *ABC* III we use only the data from the first replicate of the experiment, and for *ABC* IV only the data from the second replicate.

6.33 Mixed Series. 3×2^2 factorial, blocks of 6 units. Only the balanced plan (6.9) should be used; this may be repeated if extra replications are wanted. Thus the number of replicates must be 3, or some multiple of 3. Although the *BC* interaction is partially confounded, the relative information is high (8/9).

The statistical analysis is described, with a numerical example, in reference (6.1), p. 58.

$3^2 \times 2$ factorial, blocks of 6 units. In plan 6.11, *AB* and *ABC* are partially confounded. Either 2 or 4 replicates may be used, though 4 are required for complete balance. A numerical example for 4 replicates was given in section 6.19.

With 2 replicates (say I and II in plan 6.11) the *I* and *R* components are partially confounded, while the *J* and *S* components are unconfounded. The *I* and *R* totals are adjusted by the same formulae as in table 6.16, (iii) and (iv). The *J* and *S* totals need no adjustment. The interaction s.s. in the analysis of variance are computed as follows.

$$\begin{aligned} AB(I) &= \frac{1}{36}[(2I_1')^2 + (2I_2')^2 + (2I_3')^2] - \frac{1}{108}[2I_1' + 2I_2' + 2I_3']^2 \\ AB(J) &= \frac{1}{12}[J_1^2 + J_2^2 + J_3^2] - \frac{1}{36}[J_1 + J_2 + J_3]^2 \\ ABC(R) &= \frac{1}{12}[(2R_1')^2 + (2R_2')^2 + (2R_3')^2] \\ &\quad - \frac{1}{36}[2R_1' + 2R_2' + 2R_3']^2 \\ ABC(S) &= \frac{1}{12}[S_1^2 + S_2^2 + S_3^2] - \frac{1}{36}[S_1 + S_2 + S_3]^2 \end{aligned}$$

The *AB* two-way table and the table of individual treatment means are adjusted in the same way as with four replicates. The quantities $2I'$, $2R'$ are divided by 18 and 6, respectively, to obtain i' and r' . For i and r , we divide *I* and *R* each by 12. After taking deviations from the mean in each case, we calculate δi and δr . Thereafter the adjustment proceeds exactly as with 4 replicates, being somewhat simpler since *J* and *S* components do not enter.

3×2^3 factorial, blocks of 6 units. In plan 6.10 the interactions BC , BD , and CD , between pairs of factors that occur at 2 levels, are partially confounded with 8/9 relative information, while 5/9 relative information is retained on the interactions ABC , ABD , and ACD . Only the balanced design which requires 3 replicates is recommended.

The following computations lead to the sum of squares due to partially confounded effects. Let

$$g_1 = I_a + I_b - I_c - I_d; \quad g_1' = I_a - I_b + I_c - I_d;$$

$$g_1'' = I_a - I_b - I_c + I_d$$

where I_a is the total of block I_a , etc. Similar definitions hold for g_2, \dots, g_3'' in the other replications. Then find

$$3Q = 3[CD] + g_1 + g_2 + g_3; \quad 3Q' = 3[BD] + g_1' + g_2' + g_3';$$

$$3Q'' = 3[BC] + g_1'' + g_2'' + g_3''$$

$$3R_0 = 3[CD_{a_1}] - g_1 + g_2 + g_3; \quad 3R_1 = 3[CD_{a_1}] + g_1 + g_2 - g_3;$$

$$3R_2 = 3[CD_{a_2}] + g_1 - g_2 + g_3$$

$$3R_0' = 3[BD_{a_0}] + g_1' - g_2' + g_3'; \quad 3R_1' = 3[BD_{a_1}] - g_1' + g_2' + g_3';$$

$$3R_2' = 3[BD_{a_2}] + g_1' + g_2' - g_3'$$

$$3R_0'' = 3[BC_{a_0}] + g_1'' + g_2'' - g_3''; \quad 3R_1'' = 3[BC_{a_1}] + g_1'' - g_2'' + g_3'';$$

$$3R_2'' = 3[BC_{a_2}] - g_1'' + g_2'' + g_3''$$

The sum of squares for the two-factor interactions are as follows.

$$CD = \frac{(3Q)^2}{576}; \quad BD = \frac{(3Q')^2}{576}; \quad BC = \frac{(3Q'')^2}{576}$$

For ACD , divide the sum of squares of deviations of the quantities $3R_0, 3R_1, 3R_2$ by 120, with similar rules for ABD and ABC .

The CD , BD , and BC two-way tables must be adjusted for block effects. For the means *per unit* in the BC table, the adjustment is

$$\frac{3Q''}{192} = \frac{[BC]}{72}$$

The adjustment is added (algebraically) when b and c are both present or both absent; when b or c alone is present the adjustment is subtracted.

4^2 factorial, blocks of 4 units. In plan 6.12 each replication confounds completely a set of three of the 9 d.f. for AB . Since the relative informa-

tion on AB is $2/3$, this plan gives less precise estimates of the interactions than randomized blocks unless the reduction in block size brings a substantial decrease in the error m.s. The balanced design, which requires a multiple of 3 replicates, is preferable.

With 3 replicates the analysis of variance is as follows.

	d.f.
Replicates	2
Blocks within replicates	9
Main effects	6
Two-factor interactions	9
Error	21
	—
Total	47

To compute the sum of squares for AB , let B_{mn} be the total of the n th block in the m th replicate, and let T_{mn} be the total of all treatments that appear in this block (this will be a total over 12 units). Let $P_{mn} = T_{mn} - B_{mn}$, so that there are 12 quantities P_{mn} . Then

$$AB = \frac{1}{8}(P_{11}^2 + P_{12}^2 + \cdots + P_{34}^2) - \frac{1}{32}(G^2 + R_1^2 + R_{II}^2 + R_{III}^2)$$

where G is the grand total and the R 's are replication totals.

The estimated block mean (adjusted for treatment effects) is

$$b_{mn} = \frac{3B_{mn} - T_{mn}}{8}$$

To present a two-way table, the adjusted mean of treatment (00) is, for example,

$$(00)' = (00) - \frac{1}{8}(b_{Ib} + b_{IIa} + b_{IIIb})$$

since this treatment occurs in blocks Ib, IIa, and IIIb.

4×2^2 factorial, blocks of 8 units. In plan 6.13, one of the three components of ABC is confounded in each replicate. In other words, the difference between the treatments in block Ia and those in block Ib is one component of ABC . The plan gives $2/3$ relative information on ABC . Any number of replicates may be used, though three are needed for balanced confounding.

To obtain the sum of squares for ABC , calculate the total for each component separately, taking it only from those replicates in which it is not confounded. The divisor for the square of this total is obtained by the usual rules. Thus with 2 replicates the first component is taken from replicate II and has divisor 16, the second component is taken from replicate I and has divisor 16, while the third component comes from both replicates and has divisor 32.

If there are r replicates, the adjusted block mean is

$$b_{mn} = \frac{rB_{mn} - T_{mn}}{8(r-1)}$$

where B_{mn} is the block total and T_{mn} that of all treatments appearing in the block. With 2 replicates, the adjusted mean of treatment 111 is

$$(111)' = (111) - \frac{b_{Ia} + b_{IIb}}{2}$$

since this treatment is found in blocks Ia and IIb.

$4 \times 3 \times 2$ factorial, blocks of 12 units. The 3 d.f. for the factor A at 4 levels may be divided into the following components.

$$A' = a_3 + a_2 - a_1 - a_0$$

$$A'' = a_3 - a_2 - a_1 + a_0$$

$$A''' = a_3 - a_2 + a_1 - a_0$$

In the first 3 replicates of plan 6.14, the interactions $A'C$ and $A'BC$ are partially confounded, with 8/9 and 5/9 relative information, respectively, the interactions of A'' and A''' being unconfounded; $A''C$ and $A''BC$ are partially confounded in replicates 4 to 6 and $A'''C$ and $A'''BC$ in replicates 7 to 9. Although nine replicates are required for complete balance, the design may be used with three or six replicates. In these cases the last three or the last six replications should be taken from plan 6.14. The AC interaction suffers only a trivial loss of replication from the confounding. The statistical analysis is described in reference (6.6).

6.34 Missing Values. When factorial experiments are arranged in randomized blocks or latin squares with no confounding, the formulae previously given (sections 4.25, 4.36) are used for the estimation of missing values. With confounding, the formulae are changed, becoming in general more complicated. The different cases are discussed below.

i. *Complete confounding of the $2^3, 2^4, 2^5, 2^6, 3^3, 3^4$ factorials.* When the effects that are confounded are unimportant, we recommended that the first replication of the plan be used for all replications of the experiment, with complete confounding of these effects. In this case the estimation formula is simple. Let B be the total yield of all other units in the same block as the missing unit, and T be the total yield of all other units which have the same treatment as the missing unit. Every replication contains a block which has the same set of treatments as the block

with the missing value. Let B' be the total yield of all such blocks (including the block with the missing value).

The value for the missing unit is estimated by the formula:

$$y = \frac{rB + kT - B'}{(r-1)(k-1)}$$

where k is the number of units per block, and r is the number of replications.

It will be realized that this is the standard formula for randomized blocks (section 4.25), applied only to those blocks which contain the treatment with the missing value.

ii. *Partial confounding of the $2^3, 2^4, 2^5, 2^6, 3^3, 3^4, 4^2, 4 \times 2^2$ factorials.* The formula given later is valid when a different replication of the plan is used for each replication of the experiment. A further essential condition is that no treatment comparison be confounded in more than one replication. This condition is satisfied in all plans for the designs above, except possibly plan 6.4 (2^4 in blocks of 4) and plan 6.6 (2^6 in blocks of 8). For instance, if the first 2 replications of plan 6.6 are used, no interaction is confounded more than once; but if the first 3 replications are used, $BCDE$ is confounded twice and the formula does not apply.

The formula requires some preliminary calculations, most of which can be used in the subsequent analysis of variance. In addition to B and T , as defined in case i, we require first the total R of all other units in the same replicate as the missing unit and the grand total G of all other units in the experiment. In the other replicates, calculate the total S_b of all blocks which contain the treatment with the missing unit.

Next form the table of treatment totals. From this table compute the sum U of the observations for all other treatments which appear in the block which has the missing unit. In the other replications, calculate the corresponding sum for each block which contains the treatment with the missing unit. Let the total of these $(r-1)$ sums be V , where r is the number of replicates. Note that in the calculation of U and V the treatment with the missing value is not included.

The missing yield is estimated by the equation

$$y = \frac{kt(r-1)T + tr(r-1)B + kG + tV - krR - t(r-1)U - trS_b}{(r-1)[t(r-1)(k-1) - (t-k)]}$$

The table on p. 204 which shows the number of units whose observations are added to obtain the various totals T, B, \dots , may be helpful in checking that the totals have been correctly identified.

Consider U , for example. There are $(k-1)$ other treatments in the

block with the missing value, and each treatment total is the sum of r observations. Consequently U is a total of $r(k - 1)$ individual observations.

Total	T	B	G	V	R	U	S_b
Number of units	$(r - 1)$	$(k - 1)$	$(tr - 1)$	$r(r - 1)(k - 1)$	$(t - 1)$	$r(k - 1)$	$k(r - 1)$

iii. *Partial confounding of the 3×2^2 , 3×2^3 , $3^2 \times 2$, $4 \times 3 \times 2$ factorials.* For these designs the general method given by Yates (6.16) should be followed.

iv. *Single replication of the 2^5 , 2^6 , 3^3 , 3^4 factorials.* A missing value is estimated by minimizing the sum of squares for the interactions that are used as error.

The estimation of missing values in cases iii and iv requires a knowledge of the algebra of the analysis of variance. The expressions for the missing value are usually fairly simple in case iv but are complicated in case iii.

When the estimated value has been substituted, the analysis of variance is calculated by the appropriate method for the design. As usual, the error d.f. are reduced by one for each estimated value. Variance-ratio tests of the treatment effects are slightly disturbed. For t -tests, an approximate rule is to decrease the number of replicates ascribed to any mean by $1\frac{1}{2}$ for each estimated value that the mean contains. Suppose, for instance, that (abc) is missing in a 2^3 experiment with 3 replications. The mean of the units which receive a is taken over 12 observations, one of which is an estimated value. By the rule, the standard error of this mean is computed as $s/\sqrt{10\frac{1}{2}}$ instead of $s/\sqrt{12}$. Consequently the standard error of the mean response to A is taken as

$$\sqrt{s^2 \left(\frac{1}{10\frac{1}{2}} + \frac{1}{12} \right)}$$

instead of

$$\sqrt{\frac{2s^2}{12}}$$

6.35 Estimation of the Gain in Precision from Confounding. As mentioned previously, the advisability of confounding depends on the

amount by which the experimental error is decreased when the block size is reduced. It is therefore worth while, especially where confounding has not been previously used, to estimate the gain in precision which has been obtained in each experiment.

The technique is essentially the same for all types of design. From the results of the analysis of variance of a confounded experiment, an estimate E_r is made of the experimental error which would have been present if the experiment had been laid out in randomized complete blocks.

Single replication. Let E_b be the mean square for blocks, and E_e the mean square for error. If there is only 1 replication in the experiment, the estimate E_r is

$$E_r = \frac{n_b E_b + n_e E_e}{n_b + n_e} \quad (6.1)$$

where n_b = number of degrees of freedom for blocks.

n_e = total number of degrees of freedom *minus* n_b .

For example, with a 2^6 experiment in blocks of 8 units (single replication)

$$n_b = 7, \quad n_e = 63 - 7 = 56$$

The estimate E_r is directly comparable with the estimated error E_e . As shown in section 2.31, the comparison may be expressed in terms of the relative amounts of replication required to obtain equal accuracy with the two types of layout.

The estimate E_r is subject to the assumption that the interactions which are confounded with blocks are negligible. Since confounding is complete, no method of estimation avoids this assumption. If the confounded interactions are *not* negligible, the calculation overestimates the gain from confounding.

More than one replication. If the confounding is complete, formula 6.1 holds for E_r , subject again to the assumption that the confounded comparisons are negligible. In this case

E_b = mean square for blocks *within replications*.

E_e = error mean square.

n_b = number of degrees of freedom for blocks *within replications*.

n_e = total number of degrees of freedom *minus* number of degrees of freedom for replications *minus* n_b .

Thus with 2 replicates of a 2^6 design in blocks of 8 units, the same interactions being confounded in both replicates, we have

$$n_b = 14, \quad n_e = 127 - 1 - 14 = 112$$

With partial confounding, the same formula may be used if all confounded comparisons are assumed to be small. Evidence on the validity of this assumption is provided by the experimental results, since partially confounded comparisons can be tested for significance.

It is possible to avoid the assumption that confounded comparisons are negligible, though some extra computations are necessary. We first adjust the mean square among blocks within replicates so as to remove treatment effects. The following identity holds for the *sums of squares*.

$$\begin{aligned} \text{Blocks within replicates (adj.)} + \text{treatments (unadj.)} = \\ \text{blocks within replicates (unadj.)} + \text{treatments (adj.)} \end{aligned}$$

The two quantities on the right of the equation appear in the analysis of variance of the experimental results. Consequently the *adjusted* blocks s.s. is found from this equation by calculating the unadjusted treatments s.s. From the adjusted blocks s.s. we obtain, as shown below, a new estimate E_b' for use in formula (6.1).

These procedures may be illustrated from the results of the $3 \times 3 \times 2$ experiment in 4 replicates (section 6.19), in which AB and ABC are partially confounded. From table 6.15 the following summary of the analysis of variance is obtained.

	d.f.	s.s.	m.s.
Replications	3	3,837	1,279
Blocks within replications	8	2,836	354
Treatments	17	7,802	459
Error	43	8,909	207
Total	71	23,384	

If AB and ABC are assumed negligible, the estimate E_r is found by formula (6.1) for the case where there is more than one replication.

$$E_r = \frac{8 \times 354 + 60 \times 207}{68} = 224$$

In order to avoid the assumption, we calculate the unadjusted treatments s.s., which is found to be 8328. Since the adjusted treatments s.s. is 7802, we have

$$\text{Blocks s.s. (adj.)} = 2836 + 7802 - 8328 = 2310$$

corresponding to a mean square E_{b_a} of 289.

An unbiased estimate E_b' of the variance between blocks within replicates is

$$E_b' = \frac{4E_{b_a} - E_e}{3} = \frac{4 \times 289 - 207}{3} = 316$$

This value is inserted in formula (6.1) to give E_r' :

$$E_r' = \frac{8 \times 316 + 60 \times 207}{68} = 220$$

The estimated gain in efficiency over randomized blocks is 8% when AB and ABC are assumed negligible and 6% without this assumption. The two results agree closely, as is usually the case unless the confounded interactions are large.

The method above applies to all designs with partial confounding. For the designs discussed in this chapter, the formula for E_b' in terms of E_{ba} and E_e is

$$E_b' = \frac{rE_{ba} - E_e}{r - 1}$$

where r is the number of replicates.

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PLANS

Plan 6.1

 2^3 factorial, blocks of 4 units

Rep. I, ABC confounded

abc	ab
a	ac
b	bc
c	(1)

Plan 6.2

 2^4 factorial, blocks of 8 units

Rep. I, ABCD confounded

a	(1)
b	ab
c	ac
d	bc
abc	ad
abd	bd
acd	cd
bcd	abcd

Plan 6.3

 2^6 factorial, blocks of 16 units

Rep. I, ABCD, ABEF, CDEF confounded

a	c	ab	ac
b	d	cd	ad
acd	abc	(1)	bc
bcd	abd	abcd	bd
ce	ae	ace	abe
de	be	ade	cde
abce	acde	bce	e
abde	bcde	bde	abcde
cf	af	acf	abf
df	bf	adf	cdf
abcf	acdf	bef	f
abdf	bcdcf	bdf	abcdf
aef	cef	abef	acef
bef	def	cdef	adef
acdef	abcef	ef	bcef
bcdef	abdef	abcdef	bdef

Plan 6.4 Balanced group of sets for 2^4 factorial, blocks of 4 units

Two-factor interactions are confounded in 1 replication and three-factor interactions are confounded in 3 replications. The columns are the blocks.

Rep. I, *AB, ACD, BCD* confounded

(1)	<i>ab</i>	<i>a</i>	<i>b</i>
<i>abc</i>	<i>c</i>	<i>bc</i>	<i>ac</i>
<i>abd</i>	<i>d</i>	<i>bd</i>	<i>ad</i>
<i>cd</i>	<i>abcd</i>	<i>acd</i>	<i>bcd</i>

Rep. II, *AC, ABD, BCD*

(1)	<i>ac</i>	<i>a</i>	<i>c</i>
<i>abc</i>	<i>b</i>	<i>bc</i>	<i>ab</i>
<i>acd</i>	<i>d</i>	<i>cd</i>	<i>ad</i>
<i>bd</i>	<i>abcd</i>	<i>abd</i>	<i>bcd</i>

Rep. III, *AD, ABC, BCD*

(1)	<i>ad</i>	<i>a</i>	<i>d</i>
<i>abd</i>	<i>b</i>	<i>bd</i>	<i>ab</i>
<i>acd</i>	<i>c</i>	<i>cd</i>	<i>ac</i>
<i>bc</i>	<i>abcd</i>	<i>abc</i>	<i>bcd</i>

Rep. IV, *BC, ABD, ACD*

(1)	<i>bc</i>	<i>b</i>	<i>c</i>
<i>abc</i>	<i>a</i>	<i>ac</i>	<i>ab</i>
<i>bcd</i>	<i>d</i>	<i>cd</i>	<i>bd</i>
<i>ad</i>	<i>abcd</i>	<i>abd</i>	<i>acd</i>

Rep. V, *BD, ABC, ACD*

(1)	<i>bd</i>	<i>b</i>	<i>d</i>
<i>abd</i>	<i>a</i>	<i>ad</i>	<i>ab</i>
<i>bcd</i>	<i>c</i>	<i>cd</i>	<i>bc</i>
<i>ac</i>	<i>abcd</i>	<i>abc</i>	<i>acd</i>

Rep. VI, *CD, ABC, ABD*

(1)	<i>cd</i>	<i>c</i>	<i>d</i>
<i>acd</i>	<i>a</i>	<i>ad</i>	<i>ac</i>
<i>bcd</i>	<i>b</i>	<i>bd</i>	<i>bc</i>
<i>ab</i>	<i>abcd</i>	<i>abc</i>	<i>abd</i>

Plan 6.5 Balanced group of sets for 2^5 factorial, blocks of 8 units

Three- and four-factor interactions are confounded in 1 replication.

Rep. I, *ABC, ADE, BCDE* confounded

(1)	<i>ab</i>	<i>a</i>	<i>b</i>
<i>bc</i>	<i>ac</i>	<i>abc</i>	<i>c</i>
<i>abd</i>	<i>d</i>	<i>bd</i>	<i>ad</i>
<i>acd</i>	<i>bcd</i>	<i>be</i>	<i>abcd</i>
<i>abe</i>	<i>e</i>	<i>ce</i>	<i>ae</i>
<i>ace</i>	<i>bce</i>	<i>ade</i>	<i>abce</i>
<i>de</i>	<i>abde</i>	<i>acde</i>	<i>bde</i>
<i>bced</i>	<i>acde</i>	<i>cd</i>	<i>cde</i>

Rep. II, *ABD, BCE, ACDE*

(1)	<i>ab</i>	<i>a</i>	<i>b</i>
<i>ad</i>	<i>bd</i>	<i>d</i>	<i>abd</i>
<i>abc</i>	<i>c</i>	<i>bc</i>	<i>ac</i>
<i>bcd</i>	<i>acd</i>	<i>abcd</i>	<i>cd</i>
<i>abe</i>	<i>e</i>	<i>be</i>	<i>ae</i>
<i>bde</i>	<i>ade</i>	<i>abde</i>	<i>de</i>
<i>ce</i>	<i>abce</i>	<i>ace</i>	<i>bce</i>
<i>acde</i>	<i>bced</i>	<i>cde</i>	<i>abced</i>

Plan 6.5 (Continued)

Balanced group of sets for 2^5 factorial, blocks of 8 units

Rep. III, ACE, BCD, ABDE

(1)	ac	a	c
ae	ce	e	ace
abc	b	bc	ab
bce	abe	abce	be
acd	d	cd	ad
cde	ade	acde	de
bd	abcd	abd	bcd
abde	bde	bde	abcde

Rep. IV, ACD, BDE, ABCE

(1)	ad	a	d
ac	cd	c	acd
abd	b	bd	ab
bcd	abc	abcd	bc
ade	e	de	ae
cde	ace	acde	ce
be	abde	abe	bde
abce	bced	bce	abcde

Rep. V, ABE, CDE, ABCD

(1)	ae	a	e
ab	be	b	abe
ace	c	ce	ac
bce	abc	abce	bc
ade	d	de	ad
bde	abd	abde	bd
cd	acde	acd	cde
abcd	bced	bcd	abcde

Plan 6.6 Balanced group of sets for 2^6 factorial, blocks of 8 units

All three- and four-factor interactions are confounded in 2 replications.

Rep. I, ABC, CDE, ADF, BEF, ABDE, BCDF, ACEF confounded

abc	a	b	(1)	bc	ac	c	ab
bd	cd	abcd	acd	abd	d	ad	bcd
ae	abce	ce	bce	e	abe	be	ace
cde	bde	ade	abde	acde	bced	abcde	de
cf	bf	af	abf	acf	bef	abef	f
adf	abcdf	cdf	bcdf	df	abdf	bdf	acdf
bef	cef	abcef	acef	abef	ef	aef	bcef
abcdef	adef	bdef	def	bcedf	acdef	cdef	abdef

Rep. II, ABD, DEF, BCF, ACE, ABEF, ACDF, BCDE

abd	b	a	(1)	ad	bd	d	ab
cd	ac	bc	abc	bcd	acd	abcd	c
be	abde	de	ade	e	abe	ae	bde
ace	cde	abcde	bced	abce	ce	bce	acde
af	df	abdf	bdf	abf	f	bf	adf
bef	abcdf	cdf	acdf	cf	abcf	acf	bcd
def	acf	bef	abef	bdef	adef	abdef	ef
abcdef	bcef	acdf	cef	acdef	bcedf	cdef	abcef

Plan 6.6 (Continued)

Balanced group of sets for 2^6 factorial, blocks of 8 units

Rep. III, ABE, BDF, ACD, CEF, ADEF, BCDE, ABCF

bc	a	ac	(1)	abc	ab	b	c
acd	bd	bcd	abd	cd	d	ad	abcd
abe	ce	e	ace	be	bce	abce	ae
de	abcde	abde	bcde	ade	acde	cde	bde
af	bcf	bf	abcf	f	cf	acf	abf
bdf	acdf	adf	cdf	abdf	abcdf	bcdcf	df
cef	abef	abcef	bef	acef	aef	ef	bcef
abcdef	def	cdef	adef	bcdef	bdef	abdef	acdef

Rep. IV, ABF, CDF, ADE, BCE, ABCD, BDEF, ACEF

ac	a	b	(1)	c	abc	bc	ab
bd	bcd	acd	abcd	abd	d	ad	cd
bce	be	ae	abe	abce	ce	ace	e
ade	acde	bcde	cde	de	abde	bde	abede
abf	abcf	cf	bcf	bf	af	f	acf
cdf	df	abdf	adf	acdf	bcdcf	abcdcf	bdf
ef	cef	abcef	acef	aef	bef	abef	bcef
abcdef	abdef	def	bdef	bcdef	acdef	cdef	adef

Rep. V, ACF, BCD, ADE, BEF, ABDF, CDEF, ABCE

ab	a	bc	(1)	b	ac	c	abc
bcd	cd	abd	acd	abcd	d	ad	bd
ce	bce	ae	abce	ace	be	abe	e
ade	abde	cde	bde	de	abcde	bcde	acde
acf	abcf	f	bcf	cf	abf	bf	af
df	bdf	acdf	abdf	adf	bcdcf	abcdcf	cdf
bef	ef	abcef	aef	abef	cef	acef	bcef
abcdef	acdef	bdef	cdef	bcdef	adef	def	abdef

Rep. VI, ABC, BDE, ADF, CEF, ACDE, BCDF, ABEF

Interchange B and C in replication I

Rep. VII, ABF, DEF, BCD, ACE, ABDE, ACDF, BCEF

Interchange F and D in replication II

Rep. VIII, ABE, BDF, CDE, ACF, ADEF, ABCD, BCEF

Interchange A and E in replication III

Rep. IX, ABD, CDF, AEF, BCE, ABCF, BDEF, ACDE

Interchange F and D in replication IV

Rep. X, AEF, BDE, ACD, BCF, ABDF, CDEF, ABCE

Interchange E and C in replication V

Plan 6.7

Balanced group of sets for 3^3 factorial, blocks of 9 units

ABC confounded								
Rep. I			Rep. II					
a	b	c	a	b	c			
000	001	002	000	001	002			
011	012	010	012	010	011			
022	020	021	021	022	020			
101	102	100	102	100	101			
112	110	111	111	112	110			
120	121	122	120	121	122			
202	200	201	201	202	200			
210	211	212	210	211	212			
221	222	220	222	220	221			

Rep. III			Rep. IV					
a	b	c	a	b	c			
000	001	002	000	001	002			
011	012	010	012	010	011			
022	020	021	021	022	020			
102	100	101	101	102	100			
110	111	112	110	111	112			
121	122	120	122	120	121			
201	202	200	202	200	201			
212	210	211	211	212	210			
220	221	222	220	221	222			

Plan 6.8

Balanced group of sets for 3^4 factorial, blocks of 9 units

Three-factor interactions confounded

Rep. I, ABC IV, ABD I, ACD II, BCD III confounded

0000	0011	0022	0012	0020	0001	0021	0002	0010
1011	1022	1000	1020	1001	1012	1002	1010	1021
2022	2000	2011	2001	2012	2020	2010	2021	2002
0121	0102	0110	0100	0111	0122	0112	0120	0101
1102	1110	1121	1111	1122	1100	1120	1101	1112
2110	2121	2102	2122	2100	2111	2101	2112	2120
0212	0220	0201	0221	0202	0210	0200	0211	0222
1220	1201	1212	1202	1210	1221	1211	1222	1200
2201	2212	2220	2210	2221	2202	2222	2200	2211

*Plan 6.8 (Continued)*Balanced group of sets for 3^4 factorial, blocks of 9 units

Rep. II, ABC III, ACD IV, ABD II, BCD I

0000	0022	0011	0021	0010	0002	0012	0001	0020
1022	1011	1000	1010	1002	1021	1001	1020	1012
2011	2000	2022	2002	2021	2010	2020	2012	2001
0112	0101	0120	0100	0122	0111	0121	0110	0102
1101	1120	1112	1122	1111	1100	1110	1102	1121
2120	2112	2101	2111	2100	2122	2102	2121	2110
0221	0210	0202	0212	0201	0220	0200	0222	0211
1210	1202	1221	1201	1220	1212	1222	1211	1200
2202	2221	2210	2220	2212	2201	2211	2200	2222

Rep. III, ABC II, ACD III, ABD IV, BCD IV

0000	0021	0012	0022	0010	0001	0011	0002	0020
1021	1012	1000	1010	1001	1022	1002	1020	1011
2012	2000	2021	2001	2022	2010	2020	2011	2002
0122	0110	0101	0111	0102	0120	0100	0121	0112
1110	1101	1122	1102	1120	1111	1121	1112	1100
2101	2122	2110	2120	2111	2102	2112	2100	2121
0211	0202	0220	0200	0221	0212	0222	0210	0201
1202	1220	1211	1221	1212	1200	1210	1201	1222
2220	2211	2202	2212	2200	2221	2201	2222	2210

Rep. IV, ABC I, ACD I, ABD III, BCD II

0000	0012	0021	0011	0020	0002	0022	0001	0010
1012	1021	1000	1020	1002	1011	1001	1010	1022
2021	2000	2012	2002	2011	2020	2010	2022	2001
0111	0120	0102	0122	0101	0110	0100	0112	0121
1120	1102	1111	1101	1110	1122	1112	1121	1100
2102	2111	2120	2110	2122	2101	2121	2100	2112
0222	0201	0210	0200	0212	0221	0211	0220	0202
1201	1210	1222	1212	1221	1200	1220	1202	1211
2210	2222	2201	2221	2200	2212	2202	2211	2220

Plan 6.9 Balanced group of sets for 3×2^2 factorial, blocks of 6 units

BC, ABC confounded

Rep. I		Rep. II		Rep. III	
<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>
001	000	000	001	000	001
010	011	011	010	011	010
100	101	101	100	100	101
111	110	110	111	111	110
200	201	200	201	201	200
211	210	211	210	210	211

Plan 6.10 Balanced group of sets for 3×2^3 factorial, blocks of 6 units

BC, BD, CD
ABC, ABD, ACD confounded

Rep. I			
<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
0100	0000	0001	0010
0011	0111	0110	0101
1010	1001	1000	1100
1101	1110	1111	1011
2001	2010	2100	2000
2110	2101	2011	2111

Rep. II			
<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
0010	0001	0000	0100
0101	0110	0111	0011
1001	1010	1100	1000
1110	1101	1011	1111
2100	2000	2001	2010
2011	2111	2110	2101

Rep. III			
<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
0001	0010	0100	0000
0110	0101	0011	0111
1100	1000	1001	1010
1011	1111	1110	1101
2010	2001	2000	2100
2101	2110	2111	2011

Plan 6.11 Balanced group of sets for $3^2 \times 2$ factorial, blocks of 6 units

AB, ABC confounded

Rep. I			Rep. II		
<i>a</i>	<i>b</i>	<i>c</i>	<i>a</i>	<i>b</i>	<i>c</i>
100	200	000	200	000	100
210	010	110	010	110	210
020	120	220	120	220	020
201	001	101	101	201	001
011	111	211	211	011	111
121	221	021	021	121	221

Rep. III			Rep. IV		
<i>a</i>	<i>b</i>	<i>c</i>	<i>a</i>	<i>b</i>	<i>c</i>
100	200	000	200	000	100
010	110	210	110	210	010
220	020	120	020	120	220
201	001	101	101	201	001
111	211	011	011	111	211
021	121	221	221	021	121

Plan 6.12 Balanced group of sets for 4^2 factorial, blocks of 4 units

AB confounded

Rep. I				Rep. II				Rep. III			
<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
33	32	31	30	33	30	32	31	33	31	32	30
22	23	20	21	21	22	20	23	20	22	21	23
10	11	12	13	12	11	13	10	11	13	10	12
01	00	03	02	00	03	01	02	02	00	03	01

CONFOUNDING

Plan 6.13 Balanced group of sets for 4×2^2 factorial, blocks of 8 units

ABC confounded

Rep. I		Rep. II		Rep. III	
<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>
000	001	000	001	000	001
011	010	011	010	011	010
100	101	101	100	101	100
111	110	110	111	110	111
201	200	201	200	200	201
210	211	210	211	211	210
301	300	300	301	301	300
310	311	311	310	310	311

Plan 6.14

Balanced group of sets for $4 \times 3 \times 2$ factorial, blocks of 12 units

A'C, A'BC confounded

Rep. I		Rep. II		Rep. III	
<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>
000	001	001	000	001	000
011	010	010	011	011	010
021	020	021	020	020	021
100	101	101	100	101	100
111	110	110	111	111	110
121	120	121	120	120	121
201	200	200	201	200	201
210	211	211	210	210	211
220	221	220	221	221	220
301	300	300	301	300	301
310	311	311	310	310	311
320	321	320	321	321	320

*Plan 6.14 (Continued)*Balanced group of sets for $4 \times 3 \times 2$ factorial, blocks of 12 units $A''C, A''BC$

Rep. IV		Rep. V		Rep. VI	
<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>
001	000	000	001	000	001
010	011	011	010	010	011
020	021	020	021	021	020
100	101	101	100	101	100
111	110	110	111	111	110
121	120	121	120	120	121
200	201	201	200	201	200
211	210	210	211	211	210
221	220	221	220	220	221
301	300	300	301	300	301
310	311	311	310	310	311
320	321	320	321	321	320

$$A' = a_3 + a_2 - a_1 - a_0$$

$$A'' = a_3 - a_2 - a_1 + a_0$$

$$A''' = a_3 - a_2 + a_1 - a_0$$

 $A'''C, A'''BC$

Rep. VII		Rep. VIII		Rep. IX	
<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>	<i>a</i>	<i>b</i>
000	001	001	000	001	000
011	010	010	011	011	010
021	020	021	020	020	021
101	100	100	101	100	101
110	111	111	110	110	111
120	121	120	121	121	120
200	201	201	200	201	200
211	210	210	211	211	210
221	220	221	220	220	221
301	300	300	301	300	301
310	311	311	310	310	311
320	321	320	321	321	320

CHAPTER 7

FACTORIAL EXPERIMENTS WITH MAIN EFFECTS CONFOUNDED: SPLIT-PLOT DESIGNS

7.1 The Simple Split-plot Design

7.11 Description. In field experiments an extra factor is sometimes introduced into an experiment by dividing each plot into a number of parts. For example, if the experiment is planned originally to test a factor A with five levels, the division of each plot into halves permits the inclusion of an extra factor B at two levels. Within each plot the two levels of B are allotted at random to the two sub-plots. If the whole plots are in a randomized block design, the plan (after randomization) might appear as shown in table 7.1. It is worth noting the difference be-

TABLE 7.1 EXAMPLE OF A SPLIT-PLOT DESIGN

Rep. 1					Rep. 2					Rep. 3				
a_3	a_1	a_2	a_0	a_4	a_1	a_4	a_0	a_2	a_3	a_1	a_3	a_0	a_2	a_4
b_0	b_1	b_0	b_0	b_0	b_1	b_1	b_0	b_0	b_0	b_1	b_0	b_0	b_0	b_1
b_1	b_0	b_1	b_1	b_1	b_0	b_0	b_1	b_1	b_1	b_0	b_1	b_1	b_1	b_0

tween this arrangement and ordinary randomized blocks. In the latter the ten treatment combinations are assigned to the ten sub-plots in a replication completely at random. Here we have a more orderly assignment in which the two treatment combinations that have any given level of A always appear in the same whole-plot.

This type of arrangement is common in industrial experimentation, although the connection may not at first be realized. Frequently, one series of treatments requires rather a large bulk of experimental material, while another series can be compared with much smaller amounts. For instance, the comparison of different types of furnace for the preparation of an alloy would use much greater amounts of alloy than the comparison of different types of mould into which the alloy might be

poured. In an experiment in which both factors are to be tested, the natural procedure is to take the material prepared in any furnace, and pour some of it into each mould. That is, material prepared in one furnace at one time provides a complete replication for the comparisons among moulds, just as the plot containing any level of A provides a complete replication of the factor B . Another instance is the comparison of different machines for milking dairy cows. Each machine would necessitate rather substantial amounts of milk, whereas other comparisons, for example on the best method of pasteurizing or of cooling, could be conducted with a much smaller amount of milk per treatment. The produce from any machine could be subdivided for these subsequent tests.

We may describe this design in another way that brings out more clearly its relation to the confounded factorial designs discussed in chapter 6. If the sub-units are regarded as the experimental units, it is seen that the treatments a_0, a_1, \dots, a_4 are applied to groups or blocks of two units. Differences among these blocks are confounded with differences among the levels of A ; i.e., the main effects of A are confounded. Accordingly, the split-plot design is sometimes considered as one in which certain *main* effects are confounded, as contrasted with the designs in chapter 6, where the confounding is restricted to interactions.

7.12 Nature of the Experimental Error. In the statistical analysis, account must be taken of the fact that the observations from different sub-units in the same unit may be correlated. In field experiments this correlation is just a reflection of the fact that neighboring pieces of land tend to be similar in fertility and in other agronomic properties. In industrial experiments the same correlation may be present, because any factor that affects the whole batch of alloy prepared in one furnace at one time will tend to create similarity among smaller batches poured from it.

The mathematical analysis used to examine the effects of this correlation will be illustrated for the experiment in table 7.1. Let i refer to the replication, j to the level of A , and k to that of B : for sub-units in the same unit, the i and j subscripts will be the same. The assumption is made that a correlation ρ exists between the experimental errors e_{ijk} and e_{iju} for any two sub-units in the same unit. Sub-units in different units are assumed to be uncorrelated. Mathematically, we have

$$E(e_{ijk}e_{iju}) = \rho\sigma^2; \quad E(e_{ijk}e_{stu}) = 0$$

We now consider the effects of this model on the most important treatment comparisons. The main effects of A are calculated entirely from

unit totals or means. With two sub-units per unit, the error variance of a unit total is

$$\begin{aligned} E(e_{ij1} + e_{ij2})^2 &= E(e_{ij1}^2) + E(e_{ij2}^2) + 2E(e_{ij1}e_{ij2}) = \sigma^2 + \sigma^2 + 2\rho\sigma^2 \\ &= 2\sigma^2(1 + \rho) \end{aligned}$$

The factor 2 may be regarded as representing the effect of adding over 2 sub-units. Consequently, for the main effects of A , the appropriate error variance per sub-unit is $\sigma^2(1 + \rho)$. If there are β sub-units per unit, the corresponding variance works out at $\sigma^2[1 + (\beta - 1)\rho]$.

The main effects of B , on the other hand, are derived from the difference between the two sub-units in a unit. For this we have

$$E(e_{ij1} - e_{ij2})^2 = 2\sigma^2(1 - \rho)$$

Thus the effective variance per sub-unit applicable to the main effects of B is $\sigma^2(1 - \rho)$. This expression remains unchanged when there are β sub-units per unit. This variance also applies to any component of the AB interaction, since such components involve comparisons of $(b_1 - b_0)$ at different levels of A .

For other treatment comparisons, the basic error variance may be different from either of the two expressions above. Consider, for instance, $(a_3b_0 - a_1b_0)$, a comparison of a_3 with a_1 at the zero level of B . In any replication the 2 sub-units involved are in different units, and are therefore independent. The variance of their difference is therefore $2\sigma^2$, and the basic variance per sub-unit is σ^2 . However, the appropriate variance for all other comparisons of this type can be derived from the basic variances $\sigma^2[1 + (\beta - 1)\rho]$ and $\sigma^2(1 - \rho)$. As will be seen, the analysis of variance gives unbiased estimates of these two variances, and from these, unbiased estimates of any particular variance can be obtained.

In practice, ρ is nearly always positive. The result is that the main effects of A (the factor applied to the units) are less precisely estimated than those of B or than the AB interaction.

The analysis of variance is fairly easy. For the example in table 7.1 we first compute the 15 plot totals. Their sum of squares of deviations is partitioned in the usual way into 2 d.f. for replications, 4 for the main effects of A , and 8 for the experimental error applicable to a whole-plot. The mean square for the latter is an unbiased estimate of $\sigma^2(1 + \rho)$. All computations are divided by 2 to convert them to a sub-unit basis.

Next take the difference $(b_1 - b_0)$ on every unit. The 15 differences provide 1 d.f. which represents the main effect of B , 4 d.f. which represent the AB interactions, and the remaining 10 d.f. whose mean square gives an unbiased estimate of the sub-plot error variance $\sigma^2(1 - \rho)$. All sums

of squares are again divided by 2. The complete separation of degrees of freedom is shown in table 7.2.

TABLE 7.2 ANALYSIS OF VARIANCE FOR THE SPLIT-PLOT EXPERIMENT IN
TABLE 7.1

	d.f.
Whole plots	
Replications	2
A	4
Whole-plot error	8
	—
Total	14
Sub-plots	
B	1
AB	4
Sub-plot error	10
	—
Grand total	29

The point to be noted is that the whole-plot error is computed entirely from plot totals, and the sub-plot error entirely from the differences between sub-plots in the same plot. It will be seen that the grand total of the degrees of freedom is 29. As might be expected, the corresponding sum of squares is the sum of squares of deviations of the 30 observations from their mean. A computation of this quantity provides a check on all sums of squares. In practice, particularly with more than 2 sub-unit treatments, we usually find the sub-plot error by subtraction.

7.13 Comparison with Randomized Blocks. The experiment in table 7.1 might be arranged in ordinary randomized blocks, with 3 blocks of 10 treatments each. In a consideration of the relative merits of the two arrangements, the following points are relevant.

1. With the split-plot design, usually the *B* and the *AB* effects are estimated more precisely than the *A* effects. Moreover, the number of degrees of freedom available for the experimental error m.s. is smaller for whole-unit comparisons than for sub-unit comparisons.

2. It can be shown that the *average* experimental error over all treatment comparisons is the same for both designs. Consequently there is no *net* gain in precision resulting from the use of the split-plot design; the increased precision on *B* and *AB* is obtained by the sacrifice of precision on *A*. For tests of significance or the construction of confidence limits the randomized block design holds a slight advantage on the average since it provides more degrees of freedom for the estimate of the single error variance that it requires. For instance, the experiment cited has

8 d.f. for the whole-unit error and 10 d.f. for the sub-unit error, whereas randomized blocks would provide 18 d.f.

3. As we have indicated, the chief practical advantage of the split-plot arrangement is that it enables factors that require relatively large amounts of material and factors that require only small amounts to be combined in the same experiment. If the experiment is planned to investigate the first type of factor, so that large amounts of material are going to be used anyway, factors of the second type can often be included at very little extra cost, and some additional information obtained very cheaply.

To summarize, the split-plot design is advantageous if the B and AB effects are of greater interest than the A effects, or if the A effects cannot be tested on small amounts of material.

Two disadvantages have been mentioned by experimenters. Sometimes the whole-unit error is much larger than the sub-unit error. It may happen that the effects of A , though large and exciting, are not significant, whereas those of B , which are too small to be of practical interest, are statistically significant. The experimenter tends to be uncomfortable in reporting results of this type. Secondly, the fact that different treatment comparisons have different basic error variances makes the analysis more complex than with randomized blocks, especially if some unusual type of comparison is being made.

When the number of replications and the experimental conditions are suitable, the whole units may be arranged in a latin square. A split-plot latin square, which eliminates the error variation arising from two types of grouping, may be preferable to randomized blocks. Summarizing 22 field experiments in latin squares where the plots were split into halves, Yates (7.1) found a substantial net increase in precision over randomized blocks. The superiority of the latin square was so pronounced that even the whole-plot comparisons would have been less precisely determined in randomized block designs. Factorial combinations that lend themselves to the use of split-plot latin squares are the 5×2 , 5×3 , 5×4 , 6×2 , 6×3 , 7×2 , 7×3 , 8×2 .

7.14 Randomization. The treatments applied to the *units* are randomized according to the instructions for the design (e.g. randomized blocks, latin square, etc.) in which the units are arranged. The treatments applied to the *sub-units* are allotted at random within each unit. A separate randomization is carried out for each unit.

7.15 Statistical Analysis. If the factor A (applied to the units) contains α levels, and the factor B (applied to the sub-units) contains β

levels, the subdivision of degrees of freedom in the analysis of variance is shown in table 7.3. The details of the analysis are illustrated in section 7.17.

TABLE 7.3 PARTITION OF DEGREES OF FREEDOM FOR A SPLIT-PLOT DESIGN

Units arranged in randomized blocks (r replicates)		Units arranged in a latin square ($r = \alpha$ replicates)	
Units	d.f.	Units	d.f.
Blocks	$(r - 1)$	Rows	$(\alpha - 1)$
A	$(\alpha - 1)$	Columns	$(\alpha - 1)$
Error (a)	$(\alpha - 1)(r - 1)$	A	$(\alpha - 1)$
Total	$(r\alpha - 1)$	Error (a)	$(\alpha - 1)(\alpha - 2)$
		Total	$(\alpha^2 - 1)$
Sub-units		Sub-units	
B	$(\beta - 1)$	B	$(\beta - 1)$
AB	$(\alpha - 1)(\beta - 1)$	AB	$(\alpha - 1)(\beta - 1)$
Error (b)	$\alpha(r - 1)(\beta - 1)$	Error (b)	$\alpha(\alpha - 1)(\beta - 1)$
Total	$r\alpha(\beta - 1)$	Total	$\alpha^2(\beta - 1)$

7.16 Standard Errors. Let E_a and E_b be the error mean squares for error (a) and error (b), respectively, on a sub-unit basis. For the treatment means, also expressed on a sub-unit basis, the standard errors shown in table 7.4 apply. The final comparison in table 7.4 contains

TABLE 7.4 STANDARD ERRORS FOR THE SPLIT-PLOT DESIGN

Treatment comparison	s.e.
Difference between two A means: e.g., $[(a_1) - (a_0)]$	$\sqrt{2E_a/r\beta}$
Difference between two B means: e.g., $[(b_1) - (b_0)]$	$\sqrt{2E_b/r\alpha}$
Difference between two B means at the same level of A : e.g., $[(a_1b_1) - (a_0b_0)]$	$\sqrt{2E_b/r}$
Difference between two A means at the same level of B : e.g., $[(a_1b_1) - (a_0b_1)]$	$\sqrt{2[(\beta - 1)E_b + E_a]/r\beta}$

both the main effect of A and the AB interaction; consequently the appropriate error is a weighted mean of E_a and E_b . This error also applies to the difference between two A means which have *different* levels of B . In such cases the ratio of the treatment difference to its standard error does not follow the t -distribution. For practical purposes the approximate rule of section 4.14 may be used, though this method gives slightly too few significant results. Let t_a , t_b be the significance levels of t corre-

sponding to the degrees of freedom in E_a and E_b , respectively. The significance level of t is taken as

$$t = \frac{(\beta - 1)E_{btb} + E_{ata}}{(\beta - 1)E_b + E_a}$$

For an application, see the next section.

7.17 Numerical Example. In an experiment on the preparation of chocolate cakes, conducted at Iowa State College (7.2), 3 recipes for preparing the batter were compared. Recipes I and II differed in that the chocolate was added at 40°C. and 60°C., respectively, while recipe III contained extra sugar. In addition, 6 different baking temperatures were tested: these ranged in 10° steps from 175° to 225°. Each time that a mix was made by any recipe, enough batter was prepared for 6 cakes, each of which was baked at a different temperature. Thus the recipes are the "whole-unit" treatments, while the baking temperatures are the "sub-unit" treatments. There were 15 replications, and it will be assumed that these were conducted serially according to a randomized blocks scheme: that is, one replication was completed before starting the next, so that differences among replicates represent time differences. (The notes suggest that this was done, though they are not quite explicit.)

A number of measurements were made on the cakes. The measurement presented here is the breaking angle. One half of a slab of cake is held fixed, while the other half is pivoted about the middle until breakage occurs. The angle through which the moving half has revolved is read on a circular scale. Since breakage is gradual, the reading tends to have a subjective element. The data are shown in table 7.5.

It is customary to compute the analysis of variance on a sub-unit basis. To avoid confusion, this should be clearly stated in the analysis of variance table itself. The calculations may be presented in three steps.

Step 1. Analyze the whole-unit totals by the method appropriate to the design in which they are arranged.

$$\text{Correction factor: } \frac{(8673)^2}{270} = 278,596$$

$$\text{Total: } \frac{(269)^2 + (260)^2 + \cdots + (155)^2}{6} - 278,596 = 11,538$$

$$\text{Replications: } \frac{(843)^2 + (820)^2 + \cdots + (479)^2}{18} - 278,596 = 10,204$$

$$\text{Recipes: } \frac{(2981)^2 + (2848)^2 + (2844)^2}{90} - 278,596 = 135$$

$$\text{Error } (a), \text{ by subtraction: } 11,538 - 10,204 - 135 = 1,199$$

TABLE 7.5 BREAKING ANGLES (DEGREES)

	Rep.	Temperature						Unit totals	
		175°	185°	195°	205°	215°	225°		
Recipe I	1	42	46	47	39	53	42	269	
	2	47	29	35	47	57	45	260	
	3	32	32	37	43	45	45	234	
	4	26	32	35	24	39	26	182	
	5	28	30	31	37	41	47	214	
	6	24	22	22	29	35	26	158	
	7	26	23	25	27	33	35	169	
	8	24	33	23	32	31	34	177	
	9	24	27	28	33	34	23	169	
	10	24	33	27	31	30	33	178	
	11	33	39	33	28	33	30	196	
	12	28	31	27	39	35	43	203	
	13	29	28	31	29	37	33	187	
	14	24	40	29	40	40	31	204	
	15	26	28	32	25	37	33	181	
Totals		437	473	462	503	580	526	2981	Rep. totals
Recipe II	1	39	46	51	49	55	42	282	843
	2	35	46	47	39	52	61	280	820
	3	34	30	42	35	42	35	218	665
	4	25	26	28	46	37	37	199	599
	5	31	30	29	35	40	36	201	583
	6	24	29	29	29	24	35	170	517
	7	22	25	26	26	29	36	164	492
	8	26	23	24	31	27	37	168	493
	9	27	26	32	28	32	33	178	492
	10	21	24	24	27	37	30	163	519
	11	20	27	33	31	28	33	172	538
	12	23	28	31	34	31	29	176	557
	13	32	35	30	27	35	30	189	574
	14	23	25	22	19	21	35	145	502
	15	21	21	28	26	27	20	143	479
Totals		403	441	476	482	517	529	2848	8673
Recipe III	1	46	44	45	46	48	63	292	
	2	43	43	43	46	47	58	280	
	3	33	24	40	37	41	38	213	
	4	38	41	38	30	36	35	218	
	5	21	25	31	35	33	23	168	
	6	24	33	30	30	37	35	189	
	7	20	21	31	24	30	33	159	
	8	24	23	21	24	21	35	148	
	9	24	18	21	26	28	28	145	
	10	26	28	27	27	35	35	178	
	11	28	25	26	25	38	28	170	
	12	24	30	28	35	33	28	178	
	13	28	29	43	28	33	37	198	
	14	19	22	27	25	25	35	153	
	15	21	28	25	25	31	25	155	
Totals		419	434	476	463	516	536	2844	
Temp. totals		1259	1348	1414	1448	1613	1591	8673	

Step 2. This concerns the sub-unit treatments. Their main effects are obtained directly.

$$\text{Temperatures: } \frac{(1259)^2 + (1348)^2 + \cdots + (1591)^2}{45} - 278,596 = 2100 \checkmark$$

The sum of squares for interactions between sub-unit and whole-unit treatments is found by subtraction. First calculate the total s.s. for the two-way table that shows both sets of treatments.

Total treatments:

$$\frac{(437)^2 + (473)^2 + \cdots + (516)^2 + (536)^2}{15} - 278,596 = 2441$$

Then,

$$\text{Recipes} \times \text{temperatures: } 2441 - 135 - 2100 = 206 \checkmark$$

Step 3. Compute the total s.s. among all sub-units.

$$\text{Total: } (42)^2 + (47)^2 + \cdots + (35)^2 + (25)^2 - 278,596 = 18,143$$

The sum of squares for error (b) is then found by subtraction in table 7.6.

TABLE 7.6 ANALYSIS OF VARIANCE OF BREAKING ANGLES (ON A SUB-UNIT BASIS) FOR THE EXPERIMENT ON CHOCOLATE CAKES

Source of variation	d.f.	s.s.	m.s.	F
Replications	14	10,204		
Recipes	2	135	67.5	1.58
Error (a)	28	1,199	42.8	
Temperatures	5	2,100✓	420.0	20.49
Linear regression	1	1,967	1,967.0	95.95
Deviations	4	133	33.2	1.62
Recipes \times temperatures	10	206✓	20.6	1.00
Error (b)	210	4,299	20.5	
Total	269	18,143✓		

The mean square for recipes, while above that for error (a), does not approach the 5% level. Temperature effects are highly significant as compared with error (b). Since there is a fairly steady increase in the breaking angle with increased temperature, the sum of squares has been divided into the component due to a linear regression on temperature and that due to deviations from the straight line. The regression coefficient amounts to an increase of 1.6° in the breaking angle for each 10° rise in baking temperature. The mean square for deviations is not sig-

nificant, though it is higher than expectation. There is no indication of any interaction. It will be observed that the error (a) m.s. is about twice as large as that for error (b).

In table 7.7 are shown the treatment means, with the principal standard errors as obtained from table 7.4.

TABLE 7.7 BREAKING ANGLE MEANS (DEGREES)

Recipe	175°	185°	195°	205°	215°	225°	Recipe means
I	29.1	31.5	30.8	33.5	38.7	35.1	33.1
II	26.9	29.4	31.7	32.1	34.5	35.3	31.6
III	27.9	28.9	31.7	30.9	34.4	35.7	31.6
Temp. means	28.0	30.0	31.4	32.2	35.9	35.4	32.1

Standard error of difference between

$$\text{Two recipe means: } \sqrt{\frac{2(42.8)}{90}} = 0.98 \text{ (28 d.f.)}$$

$$\text{Two temperature means: } \sqrt{\frac{2(20.5)}{45}} = 0.95 \text{ (210 d.f.)}$$

$$\text{Two temperature means for one recipe: } \sqrt{\frac{2(20.5)}{15}} = 1.65 \text{ (210 d.f.)}$$

$$\text{Two recipe means for a given temperature: } \sqrt{\frac{2[5(20.5) + 42.8]}{90}} = 1.80$$

The 5% levels of t are 2.05 and 1.97, respectively, for 28 and 210 d.f. Consequently, the 5% level for the last standard error above is

$$\frac{(5)(20.5)(1.97) + (42.8)(2.05)}{(5)(20.5) + (42.8)} = 1.99$$

This value always lies between the two individual t -values. In practice it need rarely be calculated.

7.18 Missing Data. The formulae for inserting estimates of missing values have been developed by Anderson (7.3). Suppose first that the observation for a single sub-unit is missing, and that this sub-unit receives the treatment combination $(a_i b_j)$. Let

U = total for the unit containing the missing observations.

$(A_i B_j)$ = total of all sub-units that receive the treatment combination $(a_i b_j)$.

(A_i) = total of all observations that receive the i th level of A .

Then the estimate to be inserted for the missing value is

$$y = \frac{rU + \beta(A_i B_j) - (A_i)}{(r-1)(\beta-1)}$$

Example. Suppose that the observation for recipe II, 195° temperature, in replication 2 is missing. (Its actual value is 47.) Then

$$U = 280 - 47 = 233; \quad (A_i B_j) = 476 - 47 = 429;$$

$$(A_i) = 2848 - 47 = 2801$$

$$y = \frac{15(233) + 6(429) - 2801}{(14)(5)} = \frac{3268}{70} = 47$$

By accident, the estimated value is the same as the actual value. If several observations are missing, repeated use is made of the formula as described in section 4.25.

In the analysis of variance 1 d.f. is subtracted from the error (*b*) d.f. for each missing observation. An unbiased estimate of E_b is obtained, but the treatment sums of squares and that for error (*a*) are biased upwards. If only a small fraction of the values is missing, it appears that these biases can be ignored; methods for obtaining unbiased estimates are given by Anderson (7.3).

Standard errors. When there are missing observations, the formulae in table 7.8 are suggested for the differences between 2 means.

TABLE 7.8 STANDARD ERRORS FOR SPLIT-PLOT EXPERIMENTS WITH MISSING DATA

Treatment comparison	s.e.
Difference between two <i>A</i> means:	$\sqrt{2[E_a + fE_b]/r\beta}$
Difference between two <i>B</i> means:	$\sqrt{2E_b[1 + (f\beta/\alpha)]/ra}$
Difference between two <i>B</i> means at the same level of <i>A</i> :	$\sqrt{2E_b[1 + (f\beta/\alpha)]/r}$
Difference between two <i>A</i> means at the same level of <i>B</i> :	$\sqrt{(2E_a/r\beta) + (2E_b)[(\beta-1) + f\beta^2]/r\beta}$

If only one value is missing in the experiment, and if a mean containing that value is compared with another mean, the factor f is $\frac{1}{2}(r-1)(\beta-1)$, and the formulae are exact, reference (7.3).

With more than 1 missing observation, the value of f depends on the locations of the missing observations. The following approximation,

developed by G. S. Watson, is correct for a number of cases but otherwise tends to be slightly too high.

$$f = \frac{k}{2(r-d)(\beta-k+c-1)}$$

When counting the values of k , c , and d , as defined below, be sure to ignore all missing observations except those that occur in the 2 means that are being compared.

k = number of missing observations.

c = number of replications which contain 1 or more missing observations.

d = number of missing observations in the sub-unit treatment ($a_i b_j$) that is most affected.

For example, suppose that in the cake experiment the values for recipe I, 175° temperature, are missing in replications 1, 2, and 3, while those for recipe I, 185° temperature, are missing in replications 3 and 4. We are comparing the means for the 2 temperatures, taken over all recipes. Then k is 5, and c is 4. The sub-unit treatment with the most missing values is recipe I, 175° temperature, which has 3 missing, so that d is 3.

The case where a complete unit is missing is discussed by Anderson (7.3).

7.2 Repeated Subdivision

7.21 Description. In order to include a new factor C at γ levels, the sub-units may each be divided into sub-sub-units. There are three experimental error variances. Errors (a) and (b) have the same functions as described previously and are calculated in the same manner, except that an extra divisor γ is introduced into all sums of squares in order to present the analysis on a sub-sub-unit basis. Error (c), which will usually be the smallest of the three, applies to the C , AC , BC , and ABC effects. The process of subdivision may be carried as far as is convenient.

The partition of degrees of freedom has already been given (table 7.3) for factors A and B . For C and its interactions, the partition is shown in table 7.9.

TABLE 7.9 ANALYSIS OF VARIANCE FOR C AND ITS INTERACTIONS

Effect	d.f.
C	$(\gamma - 1)$
AC	$(\alpha - 1)(\gamma - 1)$
BC	$(\beta - 1)(\gamma - 1)$
ABC	$(\alpha - 1)(\beta - 1)(\gamma - 1)$
Error (c)	$\alpha\beta(\gamma - 1)(\gamma - 1)$

The computations in the analysis of variance are a straightforward extension of those for the simple split-plot design. The sum of squares for AC , for instance, is obtained by calculating the total s.s. for the A by C two-way table, and subtracting the sums of squares due to A and C ; and similarly for the other factorial effects in table 7.9. Finally, the sum of squares for error (c) is found by calculating the total s.s. among sub-sub-units, and subtracting the contributions from all other items in the analysis. Worked examples are given in references (7.4) and (7.5).

7.22 Standard Errors. The formulae in table 7.4 for standard errors applicable to A and B effects remain valid apart from division by an additional factor $\sqrt{\gamma}$. For the principal comparisons which involve C effects, standard errors are shown in table 7.10. All treatment means and error mean squares are assumed to be on a sub-sub-unit basis.

TABLE 7.10 STANDARD ERRORS FOR THE SPLIT-PLOT DESIGN WITH TWO SUBDIVISIONS

Treatment comparison	s.e.
$[(c_1) - (c_0)]$	$\sqrt{\frac{2E_c}{r\alpha\beta}}$
$[(a_1c_1) - (a_1c_0)]$	$\sqrt{\frac{2E_c}{r\beta}}$
$[(a_1c_1) - (a_0c_1)]$	$\sqrt{\frac{2[(\gamma - 1)E_c + E_a]}{r\beta\gamma}}$
$[(b_1c_1) - (b_1c_0)]$	$\sqrt{\frac{2E_c}{r\alpha}}$
$[(b_1c_1) - (b_0c_1)]$	$\sqrt{\frac{2[(\gamma - 1)E_c + E_b]}{r\alpha\gamma}}$

7.3 Some Variants of the Split-plot Design

7.31 Systematic Arrangement of the Treatments Applied to the Units. The device of subdividing each unit into a number of sub-units is very flexible, and can be used with any type of design in which the units are arranged. Some of the more useful variations that are possible are described in the remainder of this chapter. The first arises when the nature of the experiment makes it necessary or desirable to have the A treatments arranged in a systematic design. This arrangement has been used in experiments where the A treatments were varieties of wheat

known to mature at different dates, and the B treatments were preparations applied to the seed before sowing. Since each variety was to be harvested when ripe, the field operations were made easier by planting the varieties in the order in which they would be ready for harvest. The field plan is illustrated in table 7.11, where a_1 is the earliest variety and a_5 the latest.

TABLE 7.11 EXAMPLE OF SYSTEMATIC ARRANGEMENT OF TREATMENTS APPLIED TO UNITS

		a_1	a_2	a_3	a_4	a_5
Rep. I		b_0	b_1	b_1	b_0	b_0
		b_1	b_0	b_0	b_1	b_1
II		b_0	b_0	b_1	b_1	b_0
		b_1	b_1	b_0	b_0	b_1
III		b_0	b_0	b_0	b_1	b_0
		b_1	b_1	b_1	b_0	b_1

The sub-unit treatments are randomized within each unit; the unit treatments, however, appear in separate strips.

This arrangement provides no valid estimate of error for the A main effects, or for comparisons such as $(a_1b_0 - a_0b_0)$ that involve A effects. Thus the "whole-plot" analysis of variance is irrelevant. In the sub-plot analysis of variance, error (b) is still valid for testing the B effects and the AB interactions. One point should be watched. A comparison such as $(a_1b_1 - a_1b_0)$ is derived entirely from the first strip in the plan. Consequently, if the sub-plot error variance differs from strip to strip, the use of the error (b) m.s. for testing this quantity is not justified; instead, it is necessary to compute a separate error using only data from the first strip.

To summarize, systematic placement of the A treatments is advisable only where it is essential in order to conduct the experiment and where no test of the A main effects is required.

7.32 Sub-unit Treatments in Strips. In a further variant the sub-unit treatments, instead of being randomized independently within each

unit, are arranged in strips across each replication. For a 5×3 design the appropriate rearrangement (after randomization) might be as shown in table 7.12.

This layout may be convenient for field experiments where it is necessary to test both factors on relatively large areas and to leave free access

TABLE 7.12 DESIGN WITH SUB-UNIT TREATMENTS IN STRIPS

Rep. I					
	a_3	a_1	a_2	a_0	a_4
b_2					
b_0					
b_1					
Rep. II					
	a_1	a_4	a_0	a_2	a_3
b_1					
b_2					
b_0					
Rep. III					
	a_1	a_3	a_0	a_2	a_4
b_2					
b_1					
b_0					

Analysis of variance

	d.f.
Replications	2
A	4
Error (a)	8
B	2
Error (b)	4
AB	8
Error (c)	16
Total	44

at both ends. Similar conditions may apply in other types of experimentation.

As with the ordinary split-plot design, the average precision over all treatment comparisons is the same as that of randomized blocks arranged on the sub-units. The present design sacrifices precision on the main effects of A and B in order to provide higher precision on the interactions, which will generally be more accurately determined than in either randomized blocks or the simple split-plot design. Since in addition the numbers of degrees of freedom for estimating the errors of the A and B

effects are likely to be small, the design is not recommended unless practical considerations necessitate its use or unless the interactions are the principal object of study. Both the *A* and *B* treatments should be randomized independently within each replication.

TABLE 7.13 LATIN SQUARE DESIGN WITH SUB-UNIT TREATMENTS IN STRIPS

	a_2	a_1	a_0	a_4	a_3
b_1					
b_0					
b_2					

	a_4	a_3	a_1	a_2	a_0
b_1					
b_0					
b_2					

	a_0	a_2	a_3	a_1	a_4
b_2					
b_1					
b_0					

	a_1	a_0	a_4	a_3	a_2
b_1					
b_2					
b_0					

	a_3	a_4	a_2	a_0	a_1
b_0					
b_2					
b_1					

Analysis of variance

	d.f.
Rows	4
Columns	4
<i>A</i>	4
Error (<i>a</i>)	12
<i>B</i>	2
Error (<i>b</i>)	8
<i>AB</i>	8
Error (<i>c</i>)	32
Total	74

In the statistical analysis, separate estimates of error are obtained for A , B , and AB . Both sets of main effects may be regarded as tested in randomized block designs, in which the sum of squares for replications (blocks) is the same for both designs. A worked example (on uniformity data) is given in reference (7.6).

Formulae for the estimated standard errors of the differences between treatment means per *sub-unit* are given in table 7.14. The mean squares E_a , E_b , and E_c are assumed to be on a sub-unit basis.

TABLE 7.14 STANDARD ERRORS WHEN SUB-UNIT TREATMENTS ARE IN STRIPS
(α = number of levels of A ; β = number of levels of B ; r = number of replicates)

Treatment comparison	s.e.
$[(a_1) - (a_0)]$	$\sqrt{2E_a/r\beta}$
$[(b_1) - (b_0)]$	$\sqrt{2E_b/r\alpha}$
$[(a_1b_1) - (a_0b_1)]$	$\sqrt{2[(\alpha - 1)E_c + E_a]/r\alpha}$
$[(a_1b_1) - (a_1b_0)]$	$\sqrt{2[(\beta - 1)E_c + E_b]/r\beta}$

One of the factors may be arranged in a latin square. With a 5×3 design in 5 replicates, a layout of this type is shown in table 7.13.

The statistical analysis is straightforward. For the B treatments, the "blocks" of the randomized blocks design are the rows of the latin square. The formulae given above for the standard errors are applicable (with r equal to α).

7.33 Sub-unit Treatments in a Latin Square. In certain cases the sub-unit treatments may be arranged in a latin square, with the prospect of a further increase in accuracy on sub-unit comparisons. There are two types of layout, of which the first will be described in this section; the second, which is used when there are only a few whole-unit treatments whose main effects are not required with high accuracy, is discussed in section 8.6.

In the first type the number of replications must be a multiple of the number of sub-unit treatments. For each whole-unit treatment, the sub-unit treatments are arranged in one or more separate latin squares. An example with 4 whole-unit treatments, 3 sub-unit treatments, and 3 replications is shown in table 7.15. The latin squares are not very clearly displayed in the plan. If we examine the three units that receive a_3 , it is seen that each b appears once in the top sub-unit, once in the middle, and once at the bottom. Thus the rows of the latin square are the positions of the sub-units within the unit. The only change from the ordinary split-plot arrangement is that differences among these rows are

eliminated from the sub-unit errors. If these differences are large, sub-unit comparisons are more precisely determined than in the ordinary design. Whole-unit comparisons are unaffected.

The designs are useful where a fairly consistent gradient is expected within each unit. Examples have been quoted previously. In experi-

TABLE 7.15 SPLIT-PLOT DESIGN WITH SUB-TREATMENTS IN A LATIN SQUARE

		a_3	a_2	a_1	a_0		
Rep. I		b_2	b_1	b_0	b_1	Analysis of variance	
		b_0	b_0	b_1	b_0		
		b_1	b_2	b_2	b_2		
		a_0	a_2	a_1	a_3		
II		b_0	b_0	b_1	b_1	Units	
		b_2	b_2	b_2	b_2	Replications	2
		b_1	b_1	b_0	b_0	A	3
						Error (a)	6
						Total	11
						Sub-units	
III		a_3	a_0	a_2	a_1	Rows	8
		b_0	b_2	b_2	b_2	B	2
		b_1	b_1	b_1	b_0	AB	6
		b_2	b_0	b_0	b_1	Error (b)	8
						Total	24

ments on plant viruses, the sub-units may be the top, middle, and lower leaves of a plant, with a regular gradient of susceptibility down the plant. The sub-unit treatments may be operations performed in succession by the same person, or drugs injected in succession to the same animal, where a time-trend is anticipated.

The number of replicates may be any multiple of the number of sub-unit treatments. With 6 replicates of the design in table 7.15, the experimental material is divided into 2 groups of 3 replicates each. If practicable, the division should be made in such a way that the within-unit gradient is constant in a given group. Since the rows are eliminated separately for each group, changes in the gradient from group to group do not increase the error.

In the randomization the first step is to randomize the *units* according

to the instructions for the design in which they are arranged. For each whole-unit treatment a latin square of the size required to accommodate the sub-treatments is then randomized as described in section 4.33. A new randomization of the latin square is taken for each successive whole-unit treatment. The first column of each square is assigned to the first replication of the experiment, and so on.

For extra replication the basic plan is repeated with a new randomization.

Since the whole-unit analysis of variance is unaffected, it is necessary to show only the subdivision of the degrees of freedom among sub-units. Table 7.16 applies to the case in which the basic plan is repeated k times.

TABLE 7.16 PARTITION OF DEGREES OF FREEDOM AMONG SUB-UNITS WHEN SUB-TREATMENTS FORM LATIN SQUARES

	d.f.
Rows	$k\alpha(\beta - 1)$
Sub-unit treatments	$(\beta - 1)$
Sub-unit \times whole-unit treatments	$(\alpha - 1)(\beta - 1)$
Error (b)	$\alpha(\beta - 1)(r - 1 - k)$
Total	$r\alpha(\beta - 1)$

Notation: α = number of whole-unit treatments.

β = number of sub-unit treatments.

r = number of replications = $k\beta$.

The number of degrees of freedom for rows should be noted. Since each latin square supplies $(\beta - 1)$ degrees of freedom for rows, every whole-unit treatment contributes $k(\beta - 1)$ degrees of freedom, so that the total is $k\alpha(\beta - 1)$ degrees of freedom. The sum of squares for rows is calculated by addition of the sums of squares for each of the $k\alpha$ latin squares. The sum of squares among the columns of the latin squares constitutes a whole-unit comparison and does not appear in the portion of the analysis shown above.

Before using this design it is advisable to verify that the number of degrees of freedom for error (b) is adequate. In the example in table 7.15 there are only 8 d.f., as against 16 with an ordinary split-plot arrangement. From section 2.31 it appears that the sub-unit error variance must be reduced by at least 15% in order to compensate for the decrease in degrees of freedom.

7.34 Split-plot Technique with Confounded Designs. The treatments applied to the units may constitute a factorial system arranged in a design that confounds certain interactions completely or partially.

Since experimenters are sometimes uncertain how to compute the analysis of variance in such cases, the following notes indicate the method.

1. The totals for each unit are analyzed by the instructions for the confounded design in which the units are arranged.

2. In computing the sums of squares for the sub-unit treatments and for all interactions between sub-unit and whole-unit treatments, the confounding is ignored. Even if a whole-unit factorial effect is completely confounded, its interactions with sub-unit treatments are unconfounded and may be tested by means of error (b).

3. The sum of squares for error (b) is found as usual by subtracting the total s.s. on the whole units plus the total s.s. for sub-unit treatments and for interactions between sub-unit and whole-unit treatments from the total s.s. for sub-units.

A difficulty arises occasionally in the presentation of tables. Suppose that a $3 \times 3 \times 2$ design is applied to the units in blocks of 6 units, with AB and ABC partially confounded, and that a factor D at 2 levels is applied to the sub-units. In the A by B two-way table, the entries must be adjusted to eliminate block effects. The problem is how to present the ABD three-way table. The observed means $(a_0b_0d_0)$ and $(a_0b_0d_1)$ will average to the *unadjusted* mean for (a_0b_0) , which contains block effects. The simplest process is to calculate the difference between the adjusted and the unadjusted means of (a_0b_0) . This difference is then applied to $(a_0b_0d_0)$ and $(a_0b_0d_1)$. Notice that this adjustment does not change the difference between these two entries and brings their mean to the correct value.

7.35 Confounding of Comparisons among Sub-unit Treatments.

When the sub-unit treatments are factorial, it may be advantageous to confound certain interactions among these factors with the units. The technique, first presented by Yates (7.7), is illustrated in table 7.17, which shows a single replication of the design. The factor A applied to the units has 2 levels, while the sub-unit treatments are the 8 combinations of factors B , C , and D at 2 levels each. The ordinary split-plot arrangement is shown in (i), while (ii) and (iii) are two alternatives for the new method.

Consider first (i) and (ii). In the ordinary arrangement the unit contains 8 sub-units, to which the 8 sub-treatments are allotted at random. With the new arrangement the sub-units are in groups of 4, chosen so that BCD is confounded with group totals. The advantage of the new layout is that the size of the whole-unit has been reduced from 8 to 4 sub-units. This will in general lead to more accurate estimates both of the A effects and of the sub-unit treatments. The only exceptions are the

high-order interactions BCD and $ABCD$. In the ordinary arrangement these interactions are derived from comparisons among sub-units in the same unit, whereas in the new design they are estimated from comparisons among the units, as the reader may verify.

Arrangement (iii) goes a step further. The a_0 and a_1 treatments which have *opposite* sets of sub-unit treatments are placed in the same sub-

TABLE 7.17 CONFOUNDING OF BCD IN A SPLIT-PLOT EXPERIMENT

i. Ordinary arrangement

a_0		a_1	
bd	d	bcd	b
c	cd	d	(1)
bcd	b	bd	bc
(1)	bc	c	cd

ii. New arrangement

a_0	a_0	a_1	a_1
cd	b	(1)	c
bd	bcd	bc	d
(1)	d	cd	bcd
bc	c	bd	b

iii. New arrangement in sub-blocks

Block Ia

a_0	a_1
cd	(1)
bd	bc
(1)	cd
bc	bd

Block Ib

a_0	a_1
b	c
bcd	d
d	bcd
c	b

block. The result is that $ABCD$ is completely confounded between sub-blocks, while BCD , like A , is orthogonal with sub-blocks. The aim is to sacrifice information on $ABCD$ in the hope of obtaining more precise estimates of A and BCD . Consequently, alternative (iii) is usually preferable to (ii).

With 5 replications of arrangement (iii), the partition of degrees of freedom is set out in table 7.18 in condensed form. There are 80 sub-units and 10 sub-blocks.

TABLE 7.18 PARTITION OF DEGREES OF FREEDOM IN A 2×2^3 DESIGN WITH SPLIT-PLOT CONFOUNDING

Units	d.f.	Sub-units	d.f.
Between sub-blocks	9	B, C, D, BC, BD, CD	6
A	1	$AB, AC, AD, ABC, ABD, ACD$	6
BCD	1	Error (b)	48
Error (a)	8		—
Total between units	19	Grand total between sub-units	79

All sums of squares are calculated in the usual way. Note that $ABCD$ does not appear, since it is a component of the sum of squares between sub-blocks.

When the sub-unit treatments form a 2^3 system, a similar technique may be used for any number of levels of the factor A . Single replications of the layouts are shown in table 7.19 for A at 2, 3, 4, 5, and 6 levels.

TABLE 7.19 SPLIT-PLOT DESIGN FOR AN $\alpha \times 2^3$ FACTORIAL SYSTEM

Arrangement of whole-units in a single replication

$\alpha = 2; (2 \times 2^3)$

I_a
 I_b

a_0	a_1'
-------	--------

a_0'	a_1
--------	-------

$\alpha = 3; (3 \times 2^3)$

I_a
 I_b

a_0	a_1	a_2
-------	-------	-------

a_0'	a_1'	a_2
--------	--------	-------

$\alpha = 4; (4 \times 2^3)$

I_a

a_0	a_1'	a_2	a_3'
-------	--------	-------	--------

I_b

a_0'	a_1	a_2'	a_3
--------	-------	--------	-------

$\alpha = 5; (5 \times 2^3)$

I_a

a_0	a_1	a_2	a_3	a_4
-------	-------	-------	-------	-------

I_b

a_0'	a_1'	a_2'	a_3'	a_4'
--------	--------	--------	--------	--------

$\alpha = 6; (6 \times 2^3)$

I_a

a_0	a_1'	a_2	a_3'	a_4	a_5'
-------	--------	-------	--------	-------	--------

I_b

a_0'	a_1	a_2'	a_3	a_4'	a_5
--------	-------	--------	-------	--------	-------

Each unit contains 4 sub-units. A prime (') implies that the sub-unit treatments are (1), bc , bd , and cd , and an unaccented a that they are b , c , d , and bcd .

The arrangements differ slightly according as the number of levels of A is even or odd. With an even level, one component of $ABCD$ is confounded between sub-blocks. For example, with A at 4 levels, this component is $(a_3 + a_1 - a_2 - a_0) BCD$. When A has an odd number of levels, we are forced to confound BCD at least partially with sub-blocks, and for simplicity it has been completely confounded.

The randomization for a single replicate is as follows. (i) Allot the sub-blocks a and b at random to the 2 halves into which the replicate has been divided. (ii) Allot the levels of A at random to units within each sub-block. (iii) Allot the appropriate set of 4 sub-treatments at random

within each unit. A separate randomization is used in each replication.

The partition of degrees of freedom in the analysis of variance is shown in table 7.20 for r replications of the basic design.

TABLE 7.20 PARTITION OF DEGREES OF FREEDOM FOR AN $\alpha \times 2^3$ SPLIT-PLOT DESIGN

	Units		Sub-units	d.f.
	α even d.f.	α odd d.f.		
Sub-blocks	$(2r - 1)$	$(2r - 1)$	Sub-treatments	6
A	$(\alpha - 1)$	$(\alpha - 1)$	A \times sub-treatments	$6(\alpha - 1)$
BCD	1		Error (b)	$6\alpha(r - 1)$
ABCD	$(\alpha - 2)$	$(\alpha - 1)$	Total	$6\alpha r$
Error (a)	$2(\alpha - 1)(r - 1)$	$2(\alpha - 1)(r - 1)$		
Total	$(2\alpha r - 1)$	$(2\alpha r - 1)$		

When α is odd, the sum of squares for BCD is omitted from the treatments; otherwise all sums of squares are obtained in the usual way. With α even, 1 d.f. is missing from $ABCD$, since it is confounded with sub-blocks. Since these designs are unlikely to be used unless $ABCD$ is small, it will usually be satisfactory to combine the $ABCD$ sum of squares with error (a). The pooled sum of squares can be found by subtraction without calculating that for $ABCD$.

A considerable number of arrangements of this type are possible, though some care is required in their construction in order to avoid confounding important comparisons. Finney (7.8) gives an account of the principles of construction and designs with worked examples, for a 4×2^2 and a 6×2^2 factorial, where the units are in a 4×4 and 6×6 latin square, respectively. Yates (7.7) gives a similar design for 6×2^3 factorial.

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CHAPTER 8

FACTORIAL EXPERIMENTS CONFOUNDED IN QUASI-LATIN SQUARES

8.1 Introduction

Yates (8.1) has constructed a number of confounded arrangements for factorial designs in latin squares. In the first group of designs (plans 8.1-8.7), some of the interactions among the factors are confounded with the rows and columns of the squares. These designs resemble ordinary latin squares in that the differences among the rows and columns of the squares are eliminated from the experimental errors of the unconfounded treatment effects; however, the designs do not possess the typical latin square property that each treatment appears once in every row and every column. Yates (8.1) has proposed the name *quasi-latin squares*.

Before one of these plans is used, it is advisable to note the interactions that are confounded and the extent to which they are confounded, particularly since more treatment effects must be confounded with latin squares than with randomized blocks. Of the eight plans shown, only one involves the confounding of two-factor interactions.

In the statistical analysis, the sums of squares for rows, columns, and all unconfounded treatment effects are calculated in the usual way. Partially confounded effects, if considered negligible, may be combined with the error, which is found by subtraction; or they may be tested for significance by calculating their sums of squares from those portions of the experiment in which the effects are unconfounded.

In a further group of designs (plans 8.8-8.14), certain treatments are applied to complete rows or columns of a latin square. These designs have been called *half-plaid* squares or *plaid* squares, from the resemblance of the plan to a Scotch plaid.

8.2 Randomization of Quasi-latin Squares

The rows and the columns of each latin square are rearranged at random. In certain plans (e.g., 8.2) each square contains several replications. The randomization must *not* be restricted in such cases with the object of keeping the replications separate.

8.3 Notes on the Plans and Statistical Analysis

2^3 factorial in a 4×4 square. There are a number of alternative designs, of which two are given in plan 8.1a and 8.1b. In the former, which requires 4 replications, $\frac{1}{2}$ relative information is retained on AB , AC , BC , and ABC , main effects being unconfounded. On account of the high degree of confounding of the two-factor interactions, this plan is not recommended unless the factors are presumed to act independently or unless the 4×4 square is considered much more accurate than alternative designs.

A numerical example illustrates the general method of analysis for all designs presented here. The data in table 8.1 represent twice the yields

TABLE 8.1 YIELDS AND STATISTICAL ANALYSIS OF A 2^3 FACTORIAL IN TWO 4×4 QUASI-LATIN SQUARES

Square I					Square II					
ABC			AC		AB	AB		BC		ABC
c	(1)	abc	ab	Totals		c	a	abc	b	
57.9	62.3	60.2	58.6	239.0	AB	62.4	64.6	68.5	63.1	258.6
a	ac	b	bc	240.1		(1)	ac	bc	ab	252.1
66.2	60.5	60.0	53.4			61.5	62.7	58.6	69.3	
abc	bc	(1)	a		BC	abc	b	(1)	ac	
64.1	61.3	63.2	59.2	247.8		66.0	56.5	63.5	67.3	
b	ab	ac	c	249.1		ab	bc	a	c	234.7
58.4	61.8	67.0	61.9		65.9	52.6	58.1	58.1		
Totals	246.6	245.9	250.4		233.1	976.0	255.8	236.4	248.7	
					d.f.	s.s.		m.s.		
Squares					1	16.10				
Rows					6	100.90		16.82		
Columns					6	112.71		18.78		
A, B, C					3	146.43				
AB					1'	18.92		18.92		
AC					1'	0.06		0.06		
BC					1'	0.33		0.33		
ABC					1'	8.27		8.27		
Error					11	101.86		9.26		
Total					31	505.58				

in bushels per acre from a soil-treatment test on soybeans, at Muscatine, Iowa. The plots were split for a varietal comparison which has been omitted. For convenience, rows and columns were rearranged so that the field plan corresponds to plan 8.1a. The factors were limestone (*A*), phosphorus (*B*), and potash (*C*); in each case the lower level implies no application of the treatment.

To obtain the sums of squares for the treatment effects, first calculate the factorial effect totals in the usual way. These are shown in the first line of table 8.2. For the partially confounded effects, these totals must

TABLE 8.2 TREATMENT EFFECTS FOR A 2^3 FACTORIAL IN 4×4 QUASI-LATIN SQUARES

	[A]	[B]	[C]	[AB]	[AC]	[BC]	[ABC]
Totals	+65.3	-18.1	-9.7	+35.7	+34.9	- 8.1	- 4.3
From rows and columns				+18.3	+35.9	-10.4	+ 7.2
Adjusted totals				+17.4	- 1.0	+ 2.3	-11.5

be adjusted. Now *AB* is completely confounded with the first 2 rows of square I and the first 2 columns of square II. Calculate *AB* as obtained from these rows and columns, i.e.,

$$239.0 - 240.1 + 255.8 - 236.4 = +18.3$$

This figure is placed in the second row of table 8.2, with corresponding figures for *AC*, *BC*, and *ABC*. The adjusted totals are found by subtracting the second from the first row. Of course, these represent just the totals for each effect as found from that part of the experiment in which the effect is unconfounded. The divisors for the squares are 32 for main effects and 16 for the other effects.

The two-way tables of means require adjustment. For the *AB* table the adjustment is

$$\frac{(+17.4)}{16} - \frac{(+35.7)}{32} = -0.03$$

This value is added (algebraically) to the (1) and (*ab*) means and subtracted from the (*a*) and (*b*) means.

Plan 8.1b completely confounds *ABC* with squares; all other effects are unconfounded. The arrangement has the peculiarity that half the treatments appear only in square I, while the other half appear only in square II. Although this layout avoids the confounding of any two-factor interactions, Yates has pointed out that any interaction of the *A*

effect with squares will appear in the analysis as a BC interaction. The results are therefore likely to be misleading if the experimental material in square I differs in responsiveness from that in square II. Where this danger seems likely, Yates suggests that the columns (or rows) of the two squares may be interlaced, so that each square comprizes a randomly chosen half of the experimental material. A numerical example of the statistical analysis is given by Yates (8.1, p. 33), using uniformity data.

2^4 factorial in an 8×8 square. Four replications are required for plan 8.2. $ABCD$ is completely confounded, and threefold replication is obtained on all three-factor interactions. The degrees of freedom subdivide as follows.

	d.f.
Rows	7
Columns	7
Treatments	14
Error	35

In the computation for ABC , ABD , ACD , or BCD , the appropriate pair of rows is deleted.

2^5 factorial in an 8×8 square. The confounding in plan 8.3 (for 2 replications) is not completely balanced. Eight of the 10 three-factor interactions and four of the 5 four-factor interactions receive only single replication. The remaining high-order interactions, ABE , CDE , $ABCD$, and $ABCDE$, are unconfounded. By a proper assignment of the letters a, b, \dots to the factors, any single three-factor interaction may be left unconfounded, though not any desired pair of interactions.

The partition of degrees of freedom is:

	d.f.
Rows	7
Columns	7
Treatments	31
Error	18

In the computation of the sum of squares for ABC , ADE , or $BCDE$, the 4 rows in the experiment which correspond to the first 4 rows of plan 8.3 are omitted. The 4 rows or columns to be omitted for any other partially confounded comparison are found similarly from plan 8.3. If the high-order interaction effects are likely to be negligible, all partially confounded effects and $ABCD$ and $ABCDE$ may be combined with the error. This procedure gives 17 d.f. for treatments and 32 for error.

2^6 factorial in an 8×8 square. In each square of plan 8.4, 8 three-factor and 6 four-factor interactions are completely confounded. If the experiment has a single replication and if only the main effects and two-factor interactions are isolated for tests of significance, the degrees of freedom are allotted as follows.

	d.f.
Rows	7
Columns	7
Main effects	6
Two-factor interactions	15
Error (estimated from high-order interactions)	28

Any unconfounded three-factor interaction may also be tested.

When there are several replications in the experiment, the squares should be chosen so that the most important three-factor interactions are free from confounding. The balanced design, which requires 5 replications, gives $3/5$ relative information on all three- and four-factor interactions.

3^3 factorial in a 9×9 square. Three or six replicates may be used (plan 8.5). Four of the 8 d.f. for ABC are completely confounded in each square. The 2 squares together give balanced confounding of ABC , with $1/2$ relative information.

When square I alone is used, the components of the analysis of variance are:

	d.f.
Rows	8
Columns	8
A, B, C	6
AB, AC, BC	12
ABC	4
Error	42

The 4 d.f. for ABC are evaluated as follows. Compute the total yields of the 3 groups of 9 treatments which are described as set II (a), (c), and (b) in square 2. The sum of squares of deviations of these 3 totals, with divisor 27, constitutes 2 of the 4 d.f. The remaining 2 are found similarly from set IV (a), (b), and (c) in square 2.

With both squares, all 8 d.f. for ABC may be computed, sets I and III being taken from the experimental data for square II, and sets II and IV from the data for square I.

3^4 factorial in a 9×9 square. Each square in plan 8.6 provides $1/2$ relative information on ABC, ABD, ACD , and BCD . For an experiment

with a single replication, the simplest subdivision of the degrees of freedom is:

	d.f.
Rows	8
Columns	8
Main effects	8
Two-factor interactions	24
Error (estimated from high-order interactions)	32

When 2 replications are available, both squares should be used. All 8 d.f. for any three-factor interaction may be tested, though each is based on only a single replication. The calculation of the sums of squares for the three-factor interactions follows the method outlined for the 3^4 factorial in blocks of 9 units (section 6.32). Plan 8.6 indicates the replication from which any component must be taken. Thus, if ABC is being calculated, we use the data from square II for ABC II and ABC IV and the data from square I for ABC I and ABC III.

4×2^2 factorial in an 8×8 square. Plan 8.7 requires 4 replications. The symbols A' , A'' , A''' refer to components of the A effects.

$$A' = a_3 + a_2 - a_1 - a_0; \quad A'' = a_3 - a_2 - a_1 + a_0;$$

$$A''' = a_3 - a_2 + a_1 - a_0$$

No two-factor interaction is confounded.

In view of the small relative information on ABC (only 1/3 on the average), the sum of squares for this term will usually be combined with the error. In this event the analysis of variance reads as follows.

	d.f.
Rows	7
Columns	7
Main effects and two-factor interactions	12
Error (by subtraction)	37

8.4 Other Quasi-latin Squares

Yates (8.1) gives experimental plans which can be used for the following factorial combinations: $3^2 \times 2$ (6×6 square), 4×2^3 (8×8 square), and 9^2 (9×9 square). Analogous to the last design are designs for 7^2 (7×7 square) and 5^2 (5×5 square). In a general discussion of methods for constructing such plans, Rao (8.3) presents designs for a 2^4 factorial in 4×4 squares.

For certain factorial combinations of treatments, the *lattice squares* (chapter 12) and the incomplete latin squares (chapter 13) confound all

treatment effects to the same extent. These arrangements may be useful in cases where the experimenter does not wish to sacrifice information on the interactions.

8.5 Estimation of the Efficiency of Quasi-latin Squares

In many cases the precision of a quasi-latin square can be compared with that of some other design which might have been used. The data from the example in table 8.1 will illustrate two types of comparison.

In this experiment a 2^3 factorial system applied to plots of soybeans was arranged in two 4×4 squares, with $1/2$ relative information on AB , AC , BC , and ABC . As an alternative, the experiment might have been laid out in randomized incomplete blocks of 4 plots each, with complete confounding of ABC (plan 6.1). Since the columns were more compact in shape than the rows, we will suppose that the columns would have been chosen as blocks.

We first estimate the error m.s. that would have been obtained with incomplete blocks. It is helpful to construct analyses of variance for the 2 designs (table 8.3) for the case where there are no treatments. The

TABLE 8.3 ANALYSES OF VARIANCE (APPLICABLE TO UNIFORMITY DATA)

Quasi-latin square			Incomplete blocks		
	d.f.	m.s.		d.f.	m.s.
Columns	6	18.78	Blocks (columns)	6	18.78
Rows	6	16.82	Intra-blocks	24	(11.15)
Intra-row-and-column	18	9.26			

mean squares to be inserted in the table are explained in the next paragraph.

With incomplete blocks, the degrees of freedom to be divided between treatments and error are the 24 d.f. *within columns* (3 from each of the 8 columns). Since the randomization ascribes to treatments a random selection from these 24 d.f., the mean square for the 24 d.f. provides an unbiased estimate of the error appropriate to the incomplete block design. The 24 d.f. subdivide into 6 representing variations among rows, and 18 representing intra-row-and-column variation. If the interactions partially confounded with the rows are regarded as negligible, the rows m.s. (16.82) from the quasi-latin square analysis provides an estimate of the 6 d.f. The quasi-latin square analysis does not supply information on the whole of the remaining 18 d.f., since in the experiment 7 of these were ascribed to treatments. By virtue of the randomization for the

quasi-latin square, however, the error m.s. (9.26), with 11 d.f., may be taken as an estimate of the mean square for the whole 18 d.f.

Accordingly, an unbiased estimate of the error m.s. for the incomplete block design is

$$\frac{18 \times 9.26 + 6 \times 16.82}{24} = 11.15$$

For the estimation of main effects, the efficiency of the quasi-latin square relative to incomplete blocks is estimated as $11.15/9.26$, or 120%. Owing to the confounding in the quasi-latin square, the relative efficiency on the interactions is only 60%.

The quasi-latin square may also be compared with ordinary randomized blocks of 8 plots, where each replication is a pair of columns. In table 8.1 the column totals are as follows.

Rep. I		Rep. II		Rep. III		Rep. IV	
246.6	245.9	250.4	233.1	255.8	236.4	248.7	257.8

With uniformity data, the randomized blocks error contains 28 d.f. (7 from each replicate). To obtain this error, the incomplete blocks error (24 d.f.) must be pooled with the 4 d.f. which measure differences between the 2 columns in each pair. Since the differences between the 2 column totals in each replication are 0.7, 17.3, 19.4, and 9.1, respectively, the *sum of squares* for the 4 d.f. is

$$\frac{(0.7)^2 + (17.3)^2 + (19.4)^2 + (9.1)^2}{8} = 94.87$$

This gives $[24(11.15) + 94.87]/28 = 12.95$ as an estimate of the randomized blocks error. The relative efficiency of the quasi-latin square is 140% on main effects and 70% on interactions.

The comparisons above require the assumption that treatment effects which are confounded with rows or columns are non-existent. By more complicated procedures, estimates can be made which avoid those assumptions. The relative efficiencies quoted above are slightly too high, because we neglected the effect of the additional error d.f. that would be available in the alternative designs (section 2.31).

8.6 Treatments Applied to Complete Rows of a Latin Square

A simple example (taken from plan 8.8) is shown below. The latin square treatments form a 2×2 factorial for factors *B* and *C*. An extra

factor A has been superimposed on the rows. The first 2 rows receive the lower level of A , and the last 2 rows the higher level.

	ABC			
(-)	b	c	(1)	bc
(-)	c	b	bc	(1)
	<hr/>			
a	(1)	bc	b	c
a	bc	(1)	c	b

A

This plan might be tried in a case where the extra factor was not suitable for application to individual units. No precise estimate of the effect of A would be expected, but some useful information on the interactions between A and the other factors might be hoped for. However, this device has one property that necessitates caution in its use. Whenever an extra factor is superimposed on the rows, certain interactions between that factor and the latin square treatments are automatically confounded with columns, as noted by Yates (8.2). In the plan above, the ABC interaction

$$(abc) + (a) + (b) + (c) - (ab) - (ac) - (bc) - (1)$$

is the difference between the first 2 columns and the last 2 columns. On the other hand, AB and AC are free of column effects. In the plans presented here, which are due to Yates (8.1), such confounding is confined as far as possible to high-order interactions. The reader who constructs his own designs should verify, before using them, what interactions have been confounded in this way.

As indicated above, these designs are useful in experiments where (i) it is not easy to apply factor A to individual units, and (ii) accurate estimates of the interactions of A , but not of its main effects, are wanted. For instance, in an experiment on an irrigated crop, we might wish to find out how the responses to plant nutrients B and C are affected by the level of the water supply. Factor A could then represent a restricted and an abundant amount of irrigation water, applied to whole rows of the square.

These *half-plaid squares* are related both to quasi-latin squares and to split-plot designs. The difference from quasi-latin squares is that in the latter the confounding is confined to interactions, whereas main effects also are confounded in half-plaid squares. The connection with split-plot designs is seen when the rows are regarded as the experimental units, and the units (or plots) as sub-units. From this point of view the

plan above may be considered as a split-plot design with the sub-units arranged in latin squares.

A moderately large number of useful half-plaid squares can be constructed; only five of the simplest are given here. In the notation employed, the factorial system for the latin square treatments is placed in parentheses (), while the number of treatments imposed on the rows is shown before the parentheses. Thus plan 8.8 shows a $2 \times (2^2)$ factorial.

As with an ordinary latin square, rows and columns must both be permuted at random. The whole-row treatments cannot be arranged in separate blocks.

The statistical analysis may be followed from the partition of the degrees of freedom, which is given in condensed form with each plan. As in the split-plot design, there are two errors. The error for the factor A applied to the rows is derived from the sum of squares among rows minus the sum of squares for A .

Most of the other treatment effects are unconfounded. Their sums of squares are calculated in the usual way and tested against the latin square error. In plans 8.9 and 8.10 certain interactions are *partially* confounded with columns; the calculations for these effects are described in the notes below. In the remainder of the plans, some interactions are *completely* confounded with rows or columns and consequently do not appear in the treatments s.s. The error s.s. is found by subtraction.

8.61 Notes on the Plans and Statistical Analysis. $2 \times (2^2)$ factorial in a 4×4 square (plan 8.8). At least two squares (4 replications) should be used.

$2 \times (3 \times 2)$ factorial in a 6×6 square (plan 8.9). Each square contains 3 replicates. BC and ABC are partially confounded with columns, the relative information being $8/9$ and $5/9$, respectively. The sums of squares for these effects are calculated by the procedure for the 3×2^2 factorial in incomplete blocks of 6 units, where the columns constitute the blocks. A numerical example is given by Yates (8.1, p. 58). Columns $Ia \dots IIIb$ are identical, respectively, with Yates' blocks $Ia \dots IIIb$. Notice that the factor applied to the rows is C .

$3 \times (3 \times 2)$ factorial in a 6×6 square (plan 8.10). Each square requires 2 replicates, while the 2 squares comprise a balanced set. Column effects are partially confounded with AB and ABC , the average relative information being $7/8$ and $5/8$, respectively. If the columns are regarded as incomplete blocks, the numerical example in section 6.19 may be followed for the computation of the sums of squares for AB and ABC .

The statistical analyses for the $2 \times (2^3)$ factorial (plan 8.11) and the $2 \times (2^4)$ factorial (plan 8.12) in 8×8 squares are straightforward. In the latter, the $BCDE$ and $ABCDE$ interactions have been combined with the error for A .

Other designs. Yates (8.1, p. 79) presents the plans for a $4 \times (2^3)$ factorial in an 8×8 square and (8.1, p. 80) for a $3 \times (3^3)$ factorial in a 9×9 square. Plans can also be constructed for a $2 \times (2^5)$ system in a 8×8 square and for a $3 \times (3^2)$ system in a 9×9 square.

8.7 Treatments Applied to Complete Rows and Columns of a Latin Square

In these designs one set of treatments A is applied to complete rows of the latin square and another set B to complete columns. Although relatively low precision is obtained on the row and column treatments, the latin square treatments and their more important interactions with the A and B treatments are subject only to the latin square error. The number of useful designs appears to be limited.

Two examples are shown in plans 8.13 and 8.14. The latin square treatments form a 2^3 and 2^4 system, respectively; the additional treatments are both at 2 levels. No two-factor interactions are confounded. The rows and columns of each square should be completely randomized.

The subdivision of degrees of freedom is indicated on each plan. Interactions which are confounded with rows and columns must be omitted from the treatments s.s.: all other treatment effects are calculated in the usual way. In plan 8.14, where the square contains a single replication, the only available estimate of error comes from the high-order interactions, unless two or more squares are used.

A 9×9 square for a 3^2 system of treatments, with row and column treatments also at 3 levels each, is given in reference (8.1), p. 81.

REFERENCES

- 8.1 YATES, F. The design and analysis of factorial experiments. *Imp. Bur. Soil Sci. Tech. Comm.* 35, 1937.
- 8.2 YATES, F. The principles of orthogonality and confounding in replicated experiments. *Jour. Agr. Sci.* 23, 108-145, 1933.
- 8.3 RAO, C. R. Confounded factorial designs in quasi-latin squares. *Sankhya* 7, 295-304, 1946.

QUASI-LATIN SQUARES

PLANS

Plan 8.1a

 2^3 factorial in two 4×4 quasi-latin squares

Square I				Square II			
ABC		AC		AB		BC	
c	(1)	abc	ab	c	a	abc	b
a	ac	b	bc	(1)	ac	bc	ab
abc	bc	(1)	a	abc	b	(1)	ac
b	ab	ac	c	ab	bc	a	c

Plan 8.1b

 2^3 factorial in two 4×4 latin squares

ABC completely confounded with squares

Square I				Square II			
(1)	bc	ac	ab	b	a	abc	c
ac	ab	(1)	bc	abc	c	a	b
bc	(1)	ab	ac	a	b	c	abc
ab	ac	bc	(1)	c	abc	b	a

Plan 8.2

 2^4 factorial in an 8×8 quasi-latin square

ABCD							
c	abcd	b	ad	a	bd	abc	cd
abd	(1)	bcd	bc	acd	ac	d	ab
d	bc	a	abcd	b	cd	abd	ac
bcd	ad	acd	bd	abc	ab	c	(1)
a	bd	c	ab	d	abcd	acd	bc
abc	ac	abd	cd	bcd	(1)	b	ad
b	ab	d	ac	c	ad	bcd	abcd
acd	cd	abc	(1)	abd	bc	a	bd

Plan 8.3

 2^5 factorial in an 8×8 quasi-latin square

ACE, BCD, ABDE				ACD, BDE, ABCE				ABC, ADE, BCDE
(1)	abe	bc	ace	abd	acd	bcd	de	
bee	ac	e	ab	bed	d	abde	acde	
cde	abcd	bde	ad	abce	ae	b	c	
bd	ade	cd	abcde	be	ce	abc	a	
abc	ce	acde	bcd	(1)	bdc	ad	abe	ABD, BCE, ACDE
acd	bcd	abce	c	ade	ab	e	bd	
abde	d	a	bc	cde	bc	acc	abcd	
ae	b	abd	de	ac	abcde	cd	bce	

Plan 8.4

 2^6 factorial in 8×8 quasi-latin squares

Square I								Rows con- founded with
Columns confounded with <i>CDE, BDF, ABE, ACF, BCEF, ABCD, ADEF</i>								
<i>abcdef</i>	<i>cef</i>	<i>bf</i>	<i>bde</i>	<i>ae</i>	<i>abc</i>	<i>adf</i>	<i>cd</i>	<i>ABC</i>
<i>cde</i>	<i>abce</i>	<i>a</i>	<i>adef</i>	<i>bef</i>	<i>cf</i>	<i>bd</i>	<i>abcdf</i>	<i>BDE</i>
<i>bdf</i>	<i>af</i>	<i>abcef</i>	<i>abcd</i>	<i>c</i>	<i>be</i>	<i>cdef</i>	<i>ade</i>	<i>ADF</i>
<i>acf</i>	<i>bcd</i>	<i>def</i>	(1)	<i>abd</i>	<i>acde</i>	<i>abef</i>	<i>bce</i>	<i>CEF</i>
<i>bc</i>	<i>acd</i>	<i>abde</i>	<i>abf</i>	<i>df</i>	<i>bcd</i>	<i>e</i>	<i>acef</i>	<i>ACDE</i>
<i>ef</i>	<i>abdef</i>	<i>ace</i>	<i>bcd</i>	<i>d</i>	<i>bcf</i>	<i>ab</i>	<i>ab</i>	<i>BCDF</i>
<i>ad</i>	<i>b</i>	<i>ce</i>	<i>cdf</i>	<i>abef</i>	<i>acf</i>	<i>abcde</i>	<i>bdef</i>	<i>ABEF</i>
<i>abe</i>	<i>de</i>	<i>bcd</i>	<i>bcef</i>	<i>acdef</i>	<i>abdf</i>	<i>ac</i>	<i>f</i>	

Square II
ABC, CDE, ADF, BEF, ABDE, BCDF, ACEF

abcdef	bf	cdf	bce	abd	ac	acf	de	BCE
ae	cd	b	def	acf	abdf	abde	bcef	CDF
cde	a	abcd	acef	df	bcf	be	abdef	ABD
adf	cef	bdef	(1)	acde	abe	abcf	bcd	AEF
abc	bde	ce	bcd	abef	acdef	ad	f	BDEF
bd	abce	ade	abf	bcd	ef	c	acdf	ACDE
bef	abcd	af	abde	bc	d	cdef	ace	ABCF
cf	adef	abcef	acd	e	bcd	bdf	ab	

Square III
ABF, CDF, ADE, BCE, ABCD, BDEF, ACEF

abcdef	be	def	bef	abd	af	ace	cd	ABD
ac	df	b	cde	acf	abde	abdef	bcef	DEF
cdf	a	abdf	acef	de	bef	bc	abcde	BCF
ade	cef	bde	(1)	acdf	abc	abef	bdf	ACE
abf	bcd	cf	bdef	abce	acdef	ad	e	ABEF
bd	abcf	acd	abe	bcd	ce	f	adef	ACDF
bce	abdef	ae	abcd	bf	d	cdef	acf	BCDE
ef	acde	abcef	adf	c	bcd	bde	ab	

Plan 8.4 (Continued) 2^6 factorial in 8×8 quasi-latin squares

Square IV

BCD, ACE, ABF, DEF, ABDE, ACDF, BCEF

<i>ab</i>	<i>acd</i>	<i>ce</i>	<i>bde</i>	<i>bcf</i>	<i>df</i>	<i>acf</i>	<i>abcdef</i>	<i>AEF</i>
<i>de</i>	<i>bce</i>	<i>abcd</i>	<i>a</i>	<i>acdef</i>	<i>abef</i>	<i>bd</i>	<i>cf</i>	<i>BDE</i>
<i>c</i>	<i>bd</i>	<i>abe</i>	<i>acde</i>	<i>af</i>	<i>abcd</i>	<i>bcef</i>	<i>def</i>	<i>ACD</i>
<i>bef</i>	<i>cdef</i>	<i>acf</i>	<i>abdf</i>	<i>abce</i>	<i>ade</i>	(1)	<i>bcd</i>	<i>BCF</i>
<i>adf</i>	<i>abcf</i>	<i>bdef</i>	<i>ef</i>	<i>cd</i>	<i>b</i>	<i>abde</i>	<i>ace</i>	<i>ABDF</i>
<i>acef</i>	<i>abdef</i>	<i>bf</i>	<i>cdf</i>	<i>e</i>	<i>bced</i>	<i>abc</i>	<i>ad</i>	<i>CDEF</i>
<i>abcde</i>	<i>ae</i>	<i>d</i>	<i>bc</i>	<i>bdef</i>	<i>cef</i>	<i>acdf</i>	<i>abf</i>	<i>ABCE</i>
<i>bcdf</i>	<i>f</i>	<i>adef</i>	<i>abcef</i>	<i>abd</i>	<i>ac</i>	<i>cde</i>	<i>be</i>	

Square V

ACF, BCD, ADE, BEF, ABDF, CDEF, ABCE

<i>abcdef</i>	<i>bd</i>	<i>bc</i>	<i>acd</i>	<i>de</i>	<i>cef</i>	<i>abe</i>	<i>af</i>	<i>ABE</i>
<i>ce</i>	<i>a</i>	<i>acdf</i>	<i>bef</i>	<i>abef</i>	<i>abcde</i>	<i>def</i>	<i>bd</i>	<i>BDF</i>
<i>bcd</i>	<i>abde</i>	<i>abcef</i>	<i>cdef</i>	<i>adf</i>	<i>ac</i>	<i>bf</i>	<i>e</i>	<i>ACD</i>
<i>bef</i>	<i>abcf</i>	<i>abd</i>	(1)	<i>ace</i>	<i>adef</i>	<i>bced</i>	<i>cdf</i>	<i>CEF</i>
<i>ade</i>	<i>cd</i>	<i>f</i>	<i>abdf</i>	<i>bced</i>	<i>be</i>	<i>acef</i>	<i>abc</i>	<i>ADEF</i>
<i>ab</i>	<i>bce</i>	<i>bdef</i>	<i>acf</i>	<i>cf</i>	<i>d</i>	<i>abcf</i>	<i>acde</i>	<i>BCDE</i>
<i>acf</i>	<i>ef</i>	<i>cde</i>	<i>abce</i>	<i>b</i>	<i>bcd</i>	<i>ad</i>	<i>abdef</i>	<i>ABCF</i>
<i>df</i>	<i>acdef</i>	<i>ae</i>	<i>bde</i>	<i>abcd</i>	<i>abf</i>	<i>c</i>	<i>bcef</i>	

Plan 8.5

 3^3 factorial in 9×9 quasi-latin squares

Square I

ABC I confounded

Set III	(a)	000	102	201	011	110	212	022	121	220	
	(c)	101	200	002	112	211	010	120	222	021	
	(b)	202	001	100	210	012	111	221	020	122	
	(a)	011	110	212	022	121	220	000	102	201	
	(c)	112	211	010	120	222	021	101	200	002	<i>ABC III</i>
	(b)	210	012	111	221	020	122	202	001	100	
	(a)	022	121	220	000	102	201	011	110	212	
	(c)	120	222	021	101	200	002	112	211	010	
	(b)	221	020	122	202	001	100	210	012	111	
Set I	(a)	(b)	(c)	(a)	(b)	(c)	(a)	(b)	(c)		

Plan 8.5 (Continued) 3^3 factorial in 9×9 quasi-latin squares

Square II

ABC II

Set IV	(a)	000	101	202	012	110	211	021	122	220
	(b)	102	200	001	111	212	010	120	221	022
	(c)	201	002	100	210	011	112	222	020	121
	(a)	012	110	211	021	122	220	000	101	202
	(b)	111	212	010	120	221	022	102	200	001
	(c)	210	011	112	222	020	121	201	002	100
	(a)	021	122	220	000	101	202	012	110	211
	(b)	120	221	022	102	200	001	111	212	010
	(c)	222	020	121	201	002	100	210	011	112
Set II	(a)	(c)	(b)	(a)	(c)	(b)	(a)	(c)	(b)	

ABC IV

Plan 8.6

3^4 factorial in 9×9 quasi-latin squares

Square I

Columns confounded with ABC II, ACD III, ABD IV,

BCD IV

0000	1011	2022	0121	1102	2110	0212	1220	2201
1021	2002	0010	1112	2120	0101	1200	2211	0222
2012	0020	1001	2100	0111	1122	2221	0202	1210
0122	1100	2111	0210	1221	2202	0001	1012	2020
1110	2121	0102	1201	2212	0220	1022	2000	0011
2101	0112	1120	2222	0200	1211	2010	0021	1002
0211	1222	2200	0002	1010	2021	0120	1101	2112
1202	2210	0221	1020	2001	0012	1111	2122	0100
2220	0201	1212	2011	0022	1000	2102	0110	1121

Rows confounded with

ABC IV
ABD I
ACD II
BCD III

Square II

Columns confounded with ABC I, ACD I, ABD III,

BCD II

0000	1022	2011	0112	1101	2120	0221	1210	2202
1012	2001	0020	1121	2110	0102	1200	2222	0211
2021	0010	1002	2100	0122	1111	2212	0201	1220
0111	1100	2122	0220	1212	2201	0002	1021	2010
1120	2112	0101	1202	2221	0210	1011	2000	0022
2102	0121	1110	2211	0200	1222	2020	0012	1001
0222	1211	2200	0001	1020	2012	0110	1102	2121
1201	2220	0212	1010	2002	0021	1122	2111	0100
2210	0202	1221	2022	0011	1000	2101	0120	1112

Rows confounded with

ABC III
ACD IV
ABD II
BCD I

QUASI-LATIN SQUARES

Plan 8.7

 4×2^2 factorial in an 8×8 quasi-latin square $A'''BC$

110	311	211	001	200	010	101	300	$A'BC$
011	100	000	210	301	201	310	111	
101	210	011	311	000	300	110	201	$A''BC$
310	001	200	010	211	111	301	100	
200	010	101	300	110	311	211	001	$A'BC$
301	201	310	111	011	100	000	210	
000	300	110	201	101	210	011	311	$A''BC$
211	111	301	100	310	001	200	010	

Plan 8.8

 $2 \times (2^2)$ factorial in a 4×4 half-plaid square ABC k squares ($2k$ replicates)

-	b	c	(1)	bc	A	Squares		$(k - 1)$
-	c	b	bc	(1)		Rows		$3k$
						A	1	
						Error	$(3k - 1)$	
						Columns		$3k$
a	(1)	bc	b	c		B, C, BC, AB, AC		5
a	bc	(1)	c	b		Error		$(9k - 5)$
						Total		$(16k - 1)$

Plan 8.9

 $2 \times (3 \times 2)$ factorial in a 6×6 half-plaid square BC, ABC partially confounded

c	ab	C	Rows	5
0	01		C	1
0	10		Error	4
0	20		Columns	5
1	00		A, B, AB, AC	7
1	11		BC	1'
1	21		ABC	2'
1	00		Error	15
1	11		Total	35
1	21			

Ia Ib IIa IIb IIIa IIIb

$2 \times (2^4)$ factorial in an 8×8 half-plaid square

$ABD, BCE, ACDE$ confounded

A	—	(1)	bd	bc	cd	ce	bcd	be	de	Rows	7
	—	bcd	c	d	bce	bde	e	cde	b	A	1
	—	ce	bcd	be	de	(1)	bd	bc	cd	Error	6
	—	bde	e	cde	b	bcd	c	d	bce	Columns	7
	BCDE									Main effects	4
ABCDE	a	bc	cd	ce	bcd	be	de	(1)	bd	Two-factor interactions	10
	a	d	b	bcd	c	cde	bce	bde	e	Three-factor	8
	a	be	de	(1)	bd	bc	cd	ce	bcd	Four-factor	3
	a	cde	bce	bde	e	d	b	bcd	c	Error	24
Total											63

 $2 \times 2 \times (2^3)$ factorial in an 8×8 plaid square

$B, ACDE, ABCDE$ confounded

	-	-	-	-	b	b	b	b	Rows		7
									A	1	
									Error	6	
	-	(1)	e	cd	cde	ce	c	de	d	Columns	7
		ce	c	de	d	cd	e	(1)	cde		
	-	cd	cde	(1)	e	de	d	ce	c	B	1
		de	d	ce	c	(1)	cde	cd	e	Error	6
A									Main effects		3
BCD	a	e	(1)	cde	cd	c	ce	d	de	Two-factor interactions	10
ABCD	a	c	de	d	ce	e	(1)	cde	cd	Three-factor	9
	a	cde	cd	e	(1)	d	de	c	ce	Four-factor	3
	a	d	ce	c	de	cde	cd	e	(1)	Error	24
									Total		63

 $2 \times 2 \times (2^4)$ factorial in an 8×8 plaid square

$B, ACD, CEF, ABCD, BCEF,$
 $ADEF, ABDEF$

[illegible]

CHAPTER 9

BALANCED AND PARTIALLY BALANCED INCOMPLETE BLOCK DESIGNS

9.1 Balanced Designs

As their name implies, these designs, introduced by Yates (9.1), are arranged in blocks or groups that are smaller than a complete replication, in order to eliminate heterogeneity to a greater extent than is possible with randomized blocks and latin squares. In the designs described in chapters 6 to 8, this reduction in the size of block was achieved by sacrificing all or part of the information on certain treatment comparisons. The present designs, on the other hand, were developed for experiments in plant breeding and selection, where it is desired to make all comparisons among pairs of treatments with equal precision. Consequently, a different method for reducing the size of block is employed.

The designs may be arranged either in randomized incomplete blocks or in quasi-latin squares. They may be *balanced* or *partially balanced*. The balanced designs will be illustrated first by simple examples of the experimental plans.

Consider the plan in table 9.1 which compares 9 treatments in incomplete blocks of 3 experimental units with 4 replications.

Every pair of treatments will be found to occur once, and only once, in the same block. For instance, treatment 1 occupies the same block

TABLE 9.1 BALANCED DESIGN FOR 9 TREATMENTS IN BLOCKS OF 3 UNITS

Block	Rep. I		Rep. II		Rep. III		Rep. IV
(1)	<u>1 2 3</u>	(4)	<u>1 4 7</u>	(7)	<u>1 8 6</u>	(10)	<u>1 5 9</u>
(2)	<u>4 5 6</u>	(5)	<u>2 5 8</u>	(8)	<u>4 2 9</u>	(11)	<u>7 2 6</u>
(3)	<u>7 8 9</u>	(6)	<u>3 6 9</u>	(9)	<u>7 5 3</u>	(12)	<u>4 8 3</u>

with treatments 2 and 3 in the first replication, with treatments 4 and 7 in the second replication, with treatments 6 and 8 in the third replication, and with treatments 5 and 9 in the fourth replication. When the results are analyzed by the method of least squares, this property, to

which the adjective *balanced* is applied, ensures that all pairs of treatments are compared with approximately the same precision, even though the differences among blocks may be large. This design belongs to the group known as *balanced lattices*, so-called because the plans are conveniently written down by drawing a square lattice, with the treatment numbers at the intersections of the lines. In the balanced lattices, the number of treatments must be an exact square while the number of units per block is the corresponding square root.

Balanced designs can be constructed for other numbers of treatments and of units per block. The plan in table 9.2 shows 7 treatments arranged in blocks of 3 units.

TABLE 9.2 BALANCED DESIGN FOR 7 TREATMENTS IN BLOCKS OF 3 UNITS

Block											
(1)	<u>1</u>	<u>2</u>	<u>4</u>	(3)	<u>3</u>	<u>4</u>	<u>6</u>	(5)	<u>1</u>	<u>5</u>	<u>6</u>
(2)	<u>2</u>	<u>3</u>	<u>5</u>	(4)	<u>4</u>	<u>5</u>	<u>7</u>	(6)	<u>2</u>	<u>6</u>	<u>7</u>

Again every pair of treatments occurs once within some block. In this case, however, the blocks cannot be grouped in separate replications, since 7 is not divisible by 3. Designs of this type are known as *balanced incomplete blocks*.

For certain numbers of treatments and units per block, both the types above can be laid out in a kind of latin square formation so as to allow the elimination of variation arising from two types of grouping. The appropriate rearrangement for the first example is shown in table 9.3.

TABLE 9.3 BALANCED DESIGN FOR 9 TREATMENTS IN 4 LATTICE SQUARES

Rep. I			Rep. II			Rep. III			Rep. IV		
Rows											
(1)	<u>1</u>	<u>2</u>	<u>3</u>	(4)	<u>1</u>	<u>4</u>	<u>7</u>	(7)	<u>1</u>	<u>6</u>	<u>8</u>
(2)	<u>4</u>	<u>5</u>	<u>6</u>	(5)	<u>2</u>	<u>5</u>	<u>8</u>	(8)	<u>9</u>	<u>2</u>	<u>4</u>
(3)	<u>7</u>	<u>8</u>	<u>9</u>	(6)	<u>3</u>	<u>6</u>	<u>9</u>	(9)	<u>5</u>	<u>7</u>	<u>3</u>

It may be verified by inspection that every pair of treatments now occurs once in the same row and also once in the same column. All comparisons between pairs of treatments are of nearly equal precision. This design is known as a *lattice square*.

The second example is rewritten somewhat differently.

Every treatment now appears in each of the 3 rows, and every pair of treatments appears together once in the same column. Since the plan

TABLE 9.4 BALANCED DESIGN FOR 7 TREATMENTS IN AN INCOMPLETE LATIN SQUARE

	Columns (Blocks)						
	(1)	(2)	(3)	(4)	(5)	(6)	(7)
Rows							
(1)	1	2	3	4	5	6	7
(2)	2	3	4	5	6	7	1
(3)	4	5	6	7	1	2	3

above could represent the first 3 rows of a 7×7 latin square, this type of design has been called an *incomplete latin square*, or alternatively a *Youden square*, after W. J. Youden, who developed these designs for greenhouse experiments.

9.2 Partially Balanced Designs

Although a balanced design can be constructed with any number of treatments and any number of units per block, the minimum number of replications is fixed by these 2 variables. In most cases this number is too large for the usual conditions of experimentation. In order to allow more freedom of choice in the number of replicates, designs which lack the complete symmetry of the balanced designs must be used.

The most useful of such designs are the *lattices*. These are constructed in the same way as balanced lattices except that there are fewer replications. The design with 2 replications (e.g., the first 2 replications in table 9.1) is called a *simple lattice*, and that with 3 replications a *triple lattice*. Similarly, with a lattice square, as in table 9.3, we may use less than the full number of replicates necessary for balance. For all these designs the number of treatments must be a perfect square.

A further set, called *cubic lattices*, is useful when the number of treatments is very large. The number of treatments is the cube of the number of units per block, so that a drastic reduction in block size is obtained. The number of replicates is 3 or some multiple of 3.

Partially balanced designs are less suitable than balanced designs in a number of ways. The statistical analysis is more complicated. It is mainly for this reason that we have not included any partially balanced designs corresponding to those in tables 9.2 and 9.4. Further, when the variation among blocks (or rows and columns) is large, some pairs of treatments are more precisely compared than others, so that several standard errors are required for tests of significance. Also, the overall

precision of the experiment is decreased by the lack of symmetry. This means that under comparable conditions a partially balanced design gives a slightly higher standard error per replication than a balanced design. These disadvantages are of minor importance in lattice designs, which will be found almost as convenient as balanced designs.

9.3 Basis of the Statistical Analysis

9.31 Analysis without Recovery of Inter-block Information. To those who have previously used only the simplest designs, one feature of the incomplete block designs will be unfamiliar. The treatment averages must be adjusted in order to secure the full accuracy available in the design. That this is necessary can be seen by examining, for example, the balanced lattice in table 9.1. In three of the 4 replications, treatments 1 and 2 are in different blocks. Consequently, if the simple averages of the 2 treatments are compared, the effects of differences among blocks are eliminated only for the first replication.

Yates developed two methods of analysis for the designs. The first is an analysis by standard least squares, reference (9.1), sometimes referred to as the "intra-block" analysis. Later he showed that comparisons among the block totals also contained information about treatment effects that can be utilized in the larger experiments. This approach is analogous to that in split-plot experiments, where it will be recalled that use is made both of comparisons within whole-units (corresponding to incomplete blocks), and of comparisons among the totals of different whole-units. In this section the original method of analysis will be outlined for balanced lattice experiments. This method has not been entirely superseded by the newer analysis, because it must be used with small experiments. The newer analysis is sketched in section 9.32.

For the original analysis the mathematical model and the assumptions are essentially the same as those for previous designs in this book, as described in section 3.2. In the balanced lattice there are k^2 treatments in blocks of k units, with $(k + 1)$ replications. Let y_{ijq} be the observation for the q th treatment, which we suppose to be in the j th block within the i th replication. The model is

$$y_{ijq} = \mu + \pi_i + \beta_{ij} + \tau_q + e_{ijq} \quad (9.1)$$

where μ , π_i , β_{ij} , and τ_q represent the effects of the mean, the replicate, the incomplete block, and the treatment, respectively, and e_{ijq} is the intra-block residual or error, assumed to be normally and independently distributed with mean zero and variance σ_e^2 . As with previous designs,

we find estimates of the parameters by minimizing

$$\sum (y_{ijq} - m - p_i - b_{ij} - t_q)^2$$

subject to the usual relations

$$\sum_i p_i = 0, \quad \sum_j b_{ij} = 0 \quad (\text{for any } i), \quad \sum_q t_q = 0 \quad (9.2)$$

The normal equation for any unknown is obtained as before by equating the observed total to the expected total over all units whose equation contains the constant. Thus, if T_q denotes the total for all $(k+1)$ units that receive the q th treatment, the normal equation for t_q is

$$T_q = (k+1)m + \sum b_{ij} + (k+1)t_q \quad (9.3)$$

the terms in the p_i having disappeared in virtue of equations (9.2), since any treatment appears in *all* replicates.

This equation does not at once give the value of t_q , since it still contains the unknowns m and b_{ij} . From the normal equation for m , it is easy to see that m is the average over the whole experiment. The normal equation for any b_{ij} is

$$B_{ij} = km + kp_i + kb_{ij} + \sum t_q \quad (9.4)$$

where B_{ij} is the observed block total, and the treatments sum is over the k treatments that are in the block. If we add these equations for all blocks that contain the q th treatment, we obtain

$$B_t = k(k+1)m + k \sum b_{ij} + \sum \sum t_q \quad (9.5)$$

where B_t is the total of all such blocks.

It is at this point that the structure of the design is important. In the blocks that contain t_q , every other treatment occurs once and only once, while t_q appears $(k+1)$ times. Hence

$$\sum \sum t_q = (k+1)t_q + \sum (\text{all other } t\text{'s}) = kt_q + \sum (\text{all } t\text{'s}) = kt_q$$

from (9.2).

This simplifies (9.5) to

$$B_t = k(k+1)m + k \sum b_{ij} + kt_q \quad (9.6)$$

We may now eliminate the unwanted term $\sum b_{ij}$ from (9.3) and (9.6), giving

$$kT_q - B_t = k^2 t_q \quad (9.7)$$

a relatively simple solution. In effect, we simply adjust the treatment total for the total of the blocks in which it appears.

Since t_q represents the *deviation* of the estimated treatment mean from the mean for all treatments, it is customary in practice to compute $(t_q + m)$, which represents the estimated mean itself. A little algebraic manipulation from (9.7) shows that

$$t_q + m = \frac{T_q}{k+1} + \frac{[kT_q - (k+1)B_t + G]}{k^2(k+1)} \quad (9.8)$$

where $G = k^2(k+1)m$ is the grand total for the whole experiment. Note that $T_q/(k+1)$ is the simple or unadjusted treatment mean. The last term on the right therefore represents the adjustment that is applied to the ordinary mean. This result will be used in a later comparison with the newer analysis.

A continuation of the analysis shows that the variance of the difference between two t_q values is $2\sigma_e^2/k$. If the experiment had been in randomized blocks, with the same number of replications, $(k+1)$, the corresponding variance would have been $2\sigma^2/(k+1)$, where σ^2 is the error variance for randomized blocks. Hence the new design, with this method of analysis, gives a more accurate experiment than randomized blocks if and only if

$$\frac{\sigma_e^2}{\sigma^2} < \frac{k}{k+1}$$

The quantity $k/(k+1)$ is called the *efficiency factor* of the design. Note that the incomplete block design would be less accurate than randomized blocks if the variation among incomplete blocks were as great as that within complete replications. With the newer method of analysis this disadvantage is largely removed.

9.32 Analysis with Recovery of Inter-block Information. This analysis rests on more difficult theory and will be outlined only in part. Formally, the mathematical model is exactly the same:

$$y_{ijq} = \mu + \pi_i + \tau_q + \beta_{ij} + e_{ijq} \quad (9.9)$$

However, the additional assumptions are made that the block effects β_{ij} are normally and independently distributed with zero means and variance σ_b^2 , and that they are independent of the e_{ijq} . It follows from the assumptions that for the difference between two observations in the *same* block, the residual variance is $2\sigma_e^2$, since the β_{ij} 's cancel, whereas for the difference between two observations in different blocks, the residual variance is $2(\sigma_e^2 + \sigma_b^2)$. The reader will note the analogy with split-plot experiments (section 7.12). With regard to the justification

for the additional assumptions, it can be said that in agricultural field experiments, for which the designs were first developed, it is usually just as reasonable to make the assumptions for the β_{ij} as for the e_{ijq} . The assumptions cannot be taken for granted, and with certain types of data we might not wish to make them.

As a result of the assumptions, observations that are in the same block are positively correlated, so that the simplest type of least squares estimation cannot be applied. Instead, we may write down the joint frequency distribution of all the observations, and use the more general method of estimation known as maximum likelihood. By a well-known device in theory, maximum likelihood estimation may be shown to be equivalent to the minimization of a weighted sum of squares, consisting of two parts. The first part is the sum of squares of deviations of the residuals from their block means; the second is the sum of squares of the residuals of the block totals. The two parts receive weights w and w'/k , respectively, where

$$w = \frac{1}{\sigma_e^2}; \quad w' = \frac{1}{\sigma_e^2 + k\sigma_b^2}$$

For the moment we suppose that w and w' are known. For the balanced lattice, the quantity to be minimized is

$$\sum w[y_{ijq} - y_{ij\cdot} - (t_q - t_{ij\cdot})]^2 + \sum \frac{w'}{k} [B_{ij} - km - kp_i - kt_{ij\cdot}]^2 \quad (9.10)$$

where $y_{ij\cdot}, t_{ij\cdot}$ denote, respectively, the observed mean of a block and the mean of the t 's that occur in the block. The first sum is over all observations, the second over all blocks.

When we differentiate with respect to a given t_q , we must note that the term $t_{ij\cdot}$ will contain t_q , with coefficient $1/k$, whenever the block in question contains the q th treatment. The derivative, when equated to zero (omitting a factor -2), gives

$$\sum w[y_{ijq} - y_{ij\cdot} - t_q + t_{ij\cdot}] + \sum \frac{w'}{k} [B_{ij} - km - kp_i - kt_{ij\cdot}] = 0 \quad (9.11)$$

The first sum is over the observations that receive the q th treatment; the second over the blocks that contain this treatment. The derivative contains an additional part arising from the terms in $t_{ij\cdot}$ in the first bracket of (9.10), but this contribution will be found to vanish.

The next step is to simplify (9.11). Both brackets involve a sum of the terms $t_{ij\cdot}$. In each case, because of the symmetry of the balanced lattice,

the sum contains all other t 's once, and t_q ($k + 1$) times. Hence

$$\sum t_{ij} = \frac{(k + 1)t_q + (\text{rest of } t\text{'s})}{k} = \frac{kt_q}{k} = t_q$$

since the total of all t 's is zero. Further, we may note that the observed block mean y_{ij} equals B_{ij}/k . Writing as before T_q for the treatment total and B_t for the total of all blocks in which the treatment appears, we have from (9.11)

$$w \left[T_q - \frac{B_t}{k} - (k + 1)t_q + t_q \right] + \frac{w'}{k} [B_t - k(k + 1)m - kt_q] = 0 \quad (9.12)$$

or

$$(kw + w')t_q + (k + 1)w'm = wT_q - (w - w')\frac{B_t}{k} \quad (9.13)$$

The constant m may be shown to be the mean for the whole experiment, or $G/k^2(k + 1)$. After some rearrangement we may express (9.13) in a form comparable with the original estimate given in (9.8), as follows.

$$t_q + m = \frac{T_q}{k + 1} + \frac{(w - w')[kT_q - (k + 1)B_t + G]}{k(k + 1)(kw + w')} \quad (9.14)$$

The adjustments are seen to be of the same form in the new as in the original analysis. The two are equal when w' is zero. This occurs when σ_b^2 is very large; that is, when differences among blocks are great. In all other cases the adjustments are smaller with the new than with the original analysis. They reduce to zero when $w' = w$, which happens only if σ_b^2 is zero, or in other words if there are no real differences among blocks. Thus the new analysis is a generalization of the original analysis, reducing to it in the extreme case where there are marked variations among incomplete blocks. At the other extreme, when the arrangement into blocks has been ineffective, the new analysis makes no adjustments for the non-existent block differences, and in fact reduces to an analysis by the method for randomized blocks.

In practice the weights w and w' are not known. Yates has shown that they can be estimated from the analysis of variance, in the same way as we estimate two separate errors in a split-plot experiment. The details will not be given. The fact that weights are estimated rather than exact introduces some additional sampling variation into the adjustments. This is unimportant in the larger experiments, but in certain of the smaller experiments the weights cannot be estimated accurately and the original analysis is recommended. For other accounts of the

analysis by different approaches, see references (9.2), (9.3), (9.4), and (9.5).

9.4 Comparison of Incomplete Block and Randomized Block Designs

So far as the experimental operations are concerned, incomplete block designs are no more difficult than randomized blocks. Some extra planning is involved in drawing up and randomizing the experimental plan, especially if care is taken to make the best possible grouping of the experimental units. According to the computing equipment and experience available, the time required for the statistical analysis may exceed that for randomized blocks by 20 to 150%.

The gain in accuracy over randomized blocks depends on the type of experimental material and may be expected to increase as the number of treatments is increased. Most of the incomplete block designs cover the range from 6 to 200 treatments, while the cubic lattices extend this range to 1000 treatments without requiring a large incomplete block. The number of treatments for which a substantial increase in accuracy is attained must be determined by experience. If the experimental material is highly variable and yet lends itself to the formation of small groups which are homogeneous, the designs may be advantageous even with small numbers of treatments. From the results of varietal trials a number of comparisons with randomized blocks have been made: see, for example, references (9.6), (9.7), and (9.8). The experiments varied in size from the 3×3 to the 11×11 , and included both lattices and lattice squares. They indicated an average gain in accuracy of the order of 25%. This means that 4 replications of an incomplete block design were about as accurate as 5 replications of randomized blocks.

The ease with which the number of replications can be increased is also a factor. The object in the new designs is to obtain the most accurate comparisons that are possible from a given number of experimental units. Accordingly, the designs are likely to be most helpful when the amount of experimental material or considerations of cost and labor force the experiment to be smaller than is desired. Where the number of replicates can be increased without difficulty, the experimenter may prefer some extra replication of a simpler design which avoids the calculation of adjustments.

There is one important property, possessed by many of the designs, which increases their attractiveness relative to randomized blocks. As the plans show, the lattice square and the lattice designs are arranged in complete replications as well as in incomplete blocks. A few of the balanced incomplete block designs can also be grouped into replications.

Such designs may be regarded as randomized block designs which have additional restrictions within each replication. It has been shown by Yates (9.2) that *these designs can be analyzed as if they were ordinary randomized blocks*. This implies that the unadjusted treatment means give unbiased estimates of the true treatment effects, and that the F - and t -tests do not lose their validity. Of course, this analysis will in general be less accurate than the complete analysis.

Thus, if there is any criterion for forming incomplete blocks, an incomplete block design is worth a trial in preference to a randomized block design *which occupies the same set of replications*. When the data have been collected, the experimenter may choose whether to analyze them as randomized blocks or to complete the full analysis, with the adjustments for incomplete block variations. In fact, if the variation among incomplete blocks is no greater than that within blocks, the complete statistical analysis reduces automatically to that for randomized blocks, as mentioned in section 9.32.

The analysis as randomized blocks may be useful in experiments where several measurements are made on each experimental unit. In the formation of incomplete blocks the units are usually grouped with regard to the most important measurement. For certain other measurements from the same experiment, the grouping may be less effective; in such cases the "randomized blocks" analysis will be satisfactory, as also with measurements of subsidiary interest, where the greatest attainable accuracy is not required.

In some types of research an appreciable number of units are likely to be injured or destroyed in the course of the experiment, so that they must be omitted from the statistical analysis. With incomplete data (except where only a few units are missing) laborious computations are required to calculate the block (or row and column) adjustments. Consequently, in experiments where missing data are of frequent occurrence, incomplete block designs cannot be used to full advantage. Even in this case nothing is lost by using an incomplete block design which can be arranged in complete replicates. If when the experiment is completed it becomes evident that an appreciable number of units must be discarded, the experimenter may use the randomized blocks analysis in which the extra complication due to missing data is smaller. The same considerations apply in cases where certain treatments may have to be ignored in the final analysis.

9.5 Comparisons with Other Designs

Yates (9.1) has discussed three other designs that have been used when the number of treatments is large.

9.51 Systematic Controls. In agricultural field experiments, a control variety is sometimes placed at regular intervals over the site, the experimental varieties being arranged in complete blocks. From the yields of the control plots a fertility index may be calculated for every plot in the experiment. These indices are used to adjust the experimental plot yields for local variations in fertility.

The method of systematic controls is very flexible, since it can be used with any number of treatments and any number of replicates. Little evidence is available about the increase in accuracy obtained from the controls, though it seems probable that the increase is seldom large if the extra space occupied by the controls is taken into account. The calculation of the best adjustments, as described by Yates (9.1), is rather tedious, but if a crude type of adjustment is made most of the potential advantages of the method may be lost.

There may be additional reasons for the presence of extra controls, e.g., for their use in an observational scoring of the experimental material. In this connection it should be noted that extra controls can be included in an incomplete block design.

9.52 Random Controls. In another method the treatments are divided into groups, the grouping remaining the same in all replicates. Each group is arranged in a separate randomized block or latin square experiment. In order to obtain comparisons between treatments that are in different groups, one or more controls are included in each group and randomized along with the other treatments. Before comparing treatments that are in different groups, we subtract from each treatment mean the mean of the controls that are in the same group. Thus the controls serve to correct for differences in the fertility of the sites on which different groups are tested.

This design allows considerable flexibility in number of treatments and amount of replication and is simple to analyze. From theoretical considerations it is likely to be inferior in accuracy to a comparable incomplete block design, if one exists. Moreover the error variance is not the same for all types of comparison. If only one control is included, the variance is twice as great for the difference between two treatments that are in different groups as for two treatments in the same group.

9.53 Split-plot Designs. This arrangement is a variant of the previous design which avoids extra controls. As in section 9.52, the treatments are first divided into a number of groups of equal sizes. Instead of testing each group on a separate site, the groups are combined into a single experiment of the split-plot type. For example, with 25 treatments and 5 replications, we might first divide the treatments into 5 groups, *A*, *B*,

C, *D*, and *E*. The groups are now regarded as whole-plot treatments, and could be arranged in a 5×5 latin square on plots 5 times as large as the basic plots. Within each group the treatments are arranged at random on the individual plots.

From the analysis for a split-plot design (section 7.15), it will be seen that we can test the difference between two treatments whether they are in the same group (i.e., whole-plot) or in different groups. Controls are no longer needed. The advantages and disadvantages of this arrangement are in general similar to those of the "random controls": the split-plot design is more accurate if the space that would have been allotted to the controls is utilized. Sometimes the split-plot design is really more appropriate than an incomplete block design because the treatments divide themselves naturally into groups, and comparisons between members of the same group are regarded as more important than comparisons between members of different groups.

9.6 Choice of Incomplete Block Design

Tables 9.5 (randomized incomplete blocks) and 9.6 (lattice squares and incomplete latin squares) form an index to the incomplete block designs in this book. Each table is arranged by number of treatments (*t*) and number of units per block (*k*) and shows the numbers of replications for which designs are available, with references to the plans. It is hoped that the tables provide a rapid means of locating a suitable design, if one has been constructed.

Where the number of treatments and the size of block are fixed in advance by the conditions of the experiment, little choice is available to the experimenter. In certain types of research, however, both the number of treatments and the size of block (or row and column) can be varied to some extent without impairing the experiment.

Where more than one design appears appropriate, a design which can be arranged in separate replicates is preferable to one which cannot, and a balanced design is preferable to a partially balanced design. These recommendations usually narrow the choice to one or two designs. For example, with 25 treatments to be compared in 8 replications, we might use (see table 9.5) either a balanced incomplete block design in blocks of 4 units, or a lattice design (partially balanced) in blocks of 5 units. The latter design, though not the former, can be laid out in separate replicates.

The relative advantages of lattice and lattice square designs can be learned only by experience. From the results of a lattice square experiment we may estimate what the standard error would have been if

TABLE 9.5 DESIGNS ARRANGED IN RANDOMIZED INCOMPLETE BLOCKS

Number of treatments, t	Number of units per block, k	Number of replications, r^*		Plan number ‡	Number of treatments, t	Number of units per block, k	Number of replications, r^*		Plan number ‡
		p.b.†	b.†				p.b.†	b.†	
4	2		3	11.1	16	4	2...	5	10.2
	3		3	11.		6		6	11.27
5	2		4	11.2		6		9	11.28
	3		6	11.		10		10	11.29
	4		4	11.	19	3		9	11.30
6	2		5	11.3		9		9	11.31
	3		5	11.4		10		10	11.32
	3		10	11.5	20	4	2, 3		10.11
	4		10	11.6	21	3		10	11.33
	5		5	11.		5		5	11.34
7	2		6	11.		7		10	11.35
	3		3	11.7	25	4		8	11.36
	4		4	11.8		5	2...	6	10.3
	6		6	11.		9		9	11.37
8	2		7	11.9	27	3	3		10.
	4		7	11.10	28	4		9	11.38
	7		7	11.		7		9	11.39
9	2		8	11.	30	5	2, 3		10.12
	3	2, 3	4	10.1	31	6		6	11.40
	4		8	11.11		10		10	11.41
	5		10	11.12	36	6	2, 3		10.7
	6		8	11.13	37	9		9	11.42
	8		8	11.	41	5		10	11.43
10	2		9	11.14	42	6	2, 3		10.13
	3		9	11.15	49	7	2...	8	10.4
	4		6	11.16	56	7	2, 3		10.14
	5		9	11.17	57	8		8	11.44
	6		9	11.18	64	4	3		10.
	9		9	11.	64	8	2...	9	10.5
11	2		10	11.	72	8	2, 3		10.15
	5		5	11.19	73	9		9	11.45
	6		6	11.20	81	9	2...	10	10.6
	10		10	11.	90	9	2, 3		10.16
12	3	2, 3		10.10	91	10		10	11.46
13	3		6	11.21	100	10	2, 3		10.8
	4		4	11.22	121	11	2...		10.(12.7)
	9		9	11.23	125	5	3		10.
15	3		7	11.24	144	12	2, 3, 4		10.9
	7		7	11.25	169	13	2...		10.(12.8)
	8		8	11.26					

* Or any multiple of this number.

† p.b. = partially balanced designs.

b. = balanced designs.

‡ References 10., 11., denote *chapters*.

TABLE 9.6 DESIGNS ARRANGED IN LATTICE SQUARES AND INCOMPLETE LATIN SQUARES

Number of treatments, t	Number of units per col. and row	Number of replications, r^*		Plan number	Number of treatments, t	Number of units per col. and row	Number of replications, r^*		Plan number
		p.b.†	b.†				p.b.†	b.†	
3	3, 5	5		13.16	11	11, 5		5	13.3
	3, 7	7		13.17		11, 6		6	13.4
	3, 8	8		13.18		11, 10		10	13.†
	3, 10	10		13.19	13	13, 4		4	13.5
4	4, 3		(3)§,6	13.†		13, 9		9	13.6
	4, 5	5		13.20	15	15, 7		7	13.7
	4, 7	7		13.21		15, 8		8	13.8
	4, 9	9		13.22	16	4, 4		5	12.2
5	5, 4		4	13.†		16, 6		6	13.9
	5, 6	6		13.23		16, 10		10	13.10
	5, 9	9		13.24	19	19, 9		9	13.11
6	6, 5		5	13.†		19, 10		10	13.12
	6, 7	7		13.25	21	21, 5		5	13.13
7	7, 3		3	13.1	25	5, 5		3	12.3
	7, 4		4	13.2	31	31, 6		6	13.4
	7, 6		6	13.†	37	37, 9		9	13.15
	7, 8	8		13.26	49	7, 7	3	4	12.4
8	7, 8		7	13.†	64	8, 8	3...	9	12.5
9	3, 3		(2)§,4	12.1	81	9, 9	3, 4	5	12.6
9	9, 8		8	13.†	121	11, 11	3...	6	12.7
10	10, 9		9	13.†	169	13, 13	3...	7	12.8

* Or any multiple of this number.

† p.b. = partially balanced designs.

b. = balanced designs.

‡ Constructed from a $t \times t$ latin square by omission of last column.

§ () Not enough degrees of freedom for error.

either of the two types of grouping had not been used. Consequently the experimenter may test additional methods of grouping by using lattice squares. If the extra grouping turns out to be ineffective, the accuracy is only slightly less than that of the corresponding lattice design. The statistical computations are, however, more laborious for lattice squares, since correction terms must be obtained for both row and column effects.

In general, the lattice squares may be expected to be successful in types of experimentation where the latin square has been found superior to randomized blocks.

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CHAPTER 10

LATTICE AND CUBIC LATTICE DESIGNS

10.1 Balanced Lattices

10.11 Description. The number of treatments must be an exact square. The size of block is the square root of this number. Incomplete blocks are combined in groups to form separate replications. The special feature of the balanced lattice, as distinguished from other lattices, is that every pair of treatments occurs once in the same incomplete block. Consequently, all pairs of treatments are compared with the same degree of precision.

The numbers of replications are rather severely restricted, as well as the numbers of treatments. The useful plans are indexed below. Bal-

BALANCED LATTICES CONTAINED IN PLANS 10.1-10.6

Number of treatments	9	16	25	49	64	81
Units per block	3	4	5	7	8	9
Replicates	4	5	6	8	9	10

anced lattice designs cannot be constructed for 36 treatments, and none has been found for 100 or 144 treatments.

10.12 Statistical Analysis. The analysis is comparatively easy. It will be illustrated by an experiment on the effects of 9 feeding treatments on the growth rates of pigs, conducted by the North Carolina Agricultural Experiment Station. The analysis has been described in more detail by Comstock *et al.* (10.1), and the nutritional information obtained from the results by Peterson *et al.* (10.2).

For pigs of a given breed, previous experience indicated that a considerable part of the variance in growth rate between animals can be ascribed to the litter. Hence the experiment was planned so that litter differences would not contribute to the intra-block error. The pigs were divided into sets of 3 litter-mates. Two sets of 3 were assigned to each block. Within a block, each treatment received one member of each set. Thus the experimental unit was composed of 2 pigs each feeding in the

same pen. The plan and growth rates are given in table 10.1; the plan has been rearranged so that it follows the same pattern as plan 10.1.

TABLE 10.1 GAINS IN WEIGHT (POUNDS PER DAY) FOR A TOTAL OF 2 PIGS

Rep. I					Rep. II				
Block	(1)	(2)	(3)	Totals	Block	(1)	(4)	(7)	Totals
(1)	(1)	(2)	(3)		(4)	(1)	(4)	(7)	
	2.20	1.84	2.18	6.22		1.19	1.20	1.15	3.54
(2)	(4)	(5)	(6)		(5)	(2)	(5)	(8)	
	2.05	0.85	1.86	4.76		2.26	1.07	1.45	4.78
(3)	(7)	(8)	(9)		(6)	(3)	(6)	(9)	
	0.73	1.60	1.76	4.09		2.12	2.03	1.63	5.78
				15.07					14.10
Rep. III					Rep. IV				
(7)	(1)	(5)	(9)		(10)	(1)	(6)	(8)	
	1.81	1.16	1.11	4.08		1.77	1.57	1.43	4.77
(8)	(2)	(6)	(7)		(11)	(2)	(4)	(9)	
	1.76	2.16	1.80	5.72		1.50	1.60	1.42	4.52
(9)	(3)	(4)	(8)		(12)	(3)	(5)	(7)	
	1.71	1.57	1.13	4.41		2.04	0.93	1.78	4.75
				14.21					14.04

Treatment totals and adjustment factors

	T	B_i	$W = (3T - 4B_i + G)$	Adjusted total $T + \mu W$	Mean per unit
1	6.97	18.61	+3.89	7.21	1.80
2	7.36	21.24	-5.46	7.02	1.76
3	8.05	21.16	-3.07	7.86	1.96
4	6.42	17.23	+7.76	6.91	1.73
5	4.01	18.37	-4.03	3.76	0.94
6	7.62	21.03	-3.84	7.38	1.84
7	5.46	18.10	+1.40	5.55	1.39
8	5.61	18.05	+2.05	5.74	1.44
9	5.92	18.47	+1.30	6.00	1.50
$G = 57.42$		172.26	0.00	57.43	

The steps in the analysis are as follows. The algebraic formulae refer to a $k \times k$ lattice in blocks of k units, with $r = (k + 1)$ replicates.

1. Calculate the block totals, the replication totals, the grand total G , and the treatment totals T , shown under the plan.

2. For each treatment, calculate the total B_i for all blocks in which

the treatment appears. For treatment 4, this is

$$B_t = 4.76 + 3.54 + 4.41 + 4.52 = 17.23$$

As a check, the total of the B_t values should be k times the total of the T values.

3. Compute the quantities

$$W = kT - (k + 1)B_t + G$$

whose sum should be exactly zero.

4. The analysis of variance is now obtained. The total s.s. and the sums of squares for replications and treatments are found in the usual way. The sum of squares for blocks within replications, adjusted for treatment effects, is

$$\frac{\sum W^2}{k^3(k+1)} = \frac{(3.89)^2 + (5.46)^2 + \dots + (1.30)^2}{108} = 1.4206$$

5. Calculate the adjustment factor

$$\mu = \frac{(E_b - E_e)}{k^2 E_b} = \frac{0.1776 - 0.0773}{9 \times 0.1776} = 0.0628$$

where E_b and E_e are the blocks and intra-block m.s., respectively. The adjusted treatment total is $(T' + \mu W)$ as shown in table 10.1. To avoid confusion, the adjusted means are shown *per unit* (total of 2 pigs) although a mean per pig would be more natural. If E_b is less than E_e , μ is taken as zero, and no adjustments are applied to the treatment totals.

TABLE 10.2 ANALYSIS OF VARIANCE FOR TOTAL GROWTH RATE OF 2 PIGS

	d.f.		s.s.	m.s.
	General	$k = 3$		
Replications	k	3	0.0774	
Treatments	$(k^2 - 1)$	8	3.2261	
Blocks (adj.)	$(k^2 - 1)$	8	1.4206	0.1776
Intra-block error	$(k - 1)(k^2 - 1)$	16	1.2368	0.0773
Total	$(k^3 + k^2 - 1)$	35	5.9609	

6. For t -tests, calculate the effective error m.s.

$$E_e' = E_e(1 + k\mu) = 0.0773(1 + 3 \times 0.0628) = 0.0919$$

The purpose of the adjustment factor is to increase E_e so as to take account of sampling errors in the block correction values μW . The ordinary rules for the calculation of t -tests may now be applied to E_e' . Thus

the variance of the difference between two adjusted treatment totals is $2rE_e'$, while that for the difference between two adjusted treatment means is $2E_e'/r$, or 0.0460. The standard error of this difference is 0.214. If means per pig were taken, the standard error would be 0.107.

7. The treatments m.s. in table 10.2 cannot be tested against the intra-block error m.s., since the former contains some block effects. For an approximate F -test, calculate the sum of squares of deviations of the adjusted treatment totals, which comes to 12.6771. This is divided by r or 4 to bring to a single-unit basis, and by $(k^2 - 1)$ or 8 to obtain the mean square, 0.3962. This is tested against the effective error m.s., 0.0919. The F -ratio, $0.3962/0.0919$, or 4.31, has 8 and 16 d.f.

8. In order to estimate the precision relative to randomized blocks, pool the mean squares for blocks (adjusted) and the intra-block error. The result is 0.1107, with 24 d.f., and is an unbiased estimate of the error variance that would have been present if the experiment had been arranged in randomized blocks. This figure is compared with the effective error m.s., 0.0919. The relative precision is $0.1107/0.0919$, or 120%. This means that 4 replications of the balanced lattice appear to have been slightly less accurate than 5 replicates of randomized blocks.

These designs may be used in factorial experiments where it is desired to avoid any sacrifice of replication on the interactions relative to that on the main effects. Some of the combinations of levels that may be tested in balanced lattices are: 3×3 , 8×2 , 4×4 , $4 \times 2 \times 2$, 2^4 , 5×5 , 7×7 , 8×8 , 4^3 , 2^6 , 9×9 , 3^4 . In analyzing a factorial experiment, first carry through the procedure described in this section. The sum of squares for deviations of the adjusted treatment totals is then divided into the sums of squares for main effects and interactions in the usual way; a divisor r must be introduced in order to convert these sums of squares to a single-unit basis. The resulting mean squares are tested against the effective error m.s., $E_e(1 + k\mu)$, which has $(k - 1)(k^2 - 1)$, or in this case, 16 d.f.

10.13 Missing Data. Since lattice designs are often used in large experiments, involving perhaps several hundred observations, it is not easy to ensure that all the observations are accurately made. Even with careful management of the experiment, there is always a chance that mistakes or accidents will affect a few of the observations. Consequently, missing data tend to be more common with lattices than with small experiments.

Methods for the analysis of the results of lattice experiments with incomplete data have been developed by Cornish (10.3). As might be expected, the computations are lengthy. They become simpler in two special cases. The first arises when the incomplete blocks are ineffective,

so that the analysis reduces to a "randomized blocks" analysis. In this event values are substituted for the missing observations by the formula which applies to randomized blocks. The second case occurs at the other extreme when the variation among blocks is so large that the inter-block information is negligible. Here the correct procedure is to insert values for the missing observations by minimizing the intra-block error m.s.

Since the general procedure must reduce to these two special procedures in the appropriate cases, it might be anticipated, as Cornish's solution shows, that both the "randomized blocks" and the "intra-block" estimates for the missing observations are required. Thus we have to find two estimates for each missing value. Similarly, two analyses of variance are necessary, one to obtain the correct value for the intra-block error m.s. and one for obtaining the mean square among blocks.

In an attempt to reduce the amount of arithmetic, some investigation has been made of the consequences of using a single estimate and a single analysis of variance. For this purpose the "intra-block" estimate seems the better of the two. It gives the correct intra-block error m.s.; the block m.s. is in general slightly too high. It provides an excellent approximation when the blocks are effective and is at its worst when blocks are ineffective; that is, when a "randomized blocks" analysis should have been used. In the latter case it still gives unbiased estimates of the treatment means, but they are not as accurate as the estimates obtained by the use of the "randomized blocks" formula. The chief defect of the "randomized blocks" formula is that when block variation is large it tends to give an overestimate of the intra-block error m.s., and sometimes the bias is substantial. For this reason the "randomized blocks" estimate is considered more hazardous for general use, despite its greater simplicity.

The procedure that is given for the incomplete block designs in this and succeeding chapters is to insert values for the missing observations by means of the "intra-block" formula. Thereafter the analysis is conducted in the usual way for complete data, except that in the analysis of variance 1 d.f. is subtracted from the intra-block error for each missing observation. This method, or the more accurate Cornish method, should be used whenever it is intended to recover inter-block information. If many observations are missing, or if it is evident on inspection that blocks are relatively ineffective, the experimenter may decide at the start to analyze the data as a randomized block experiment. In this case substitutes for the missing values should of course be obtained by the formula for randomized blocks (section 4.25).

The "intra-block" formula for the estimate of a missing observation in a $k \times k$ balanced lattice with $(k + 1)$ replications is as follows.

$$x = \frac{k^2 T + k(k + 1)B - R + G - kT_b - kB_t}{k(k - 1)^2} \quad (10.1)$$

In this formula, T , B , and R denote as usual the totals for the treatment, block, and replication which contain the missing observation, while G is the grand total. Also

T_b = total (over all replications) of all treatments that appear in the block which has the missing observation.

B_t = total of all blocks in which the treatment with the missing observation appears.

Example. Suppose that the observation 2.20 for treatment 1 in replicate 1 of table 10.1 had been missing. Form the block, replicate, treatment, and grand totals, just as in the ordinary analysis. It is helpful to insert an x for the missing observation and to include this x in all totals where it should appear. When the value of x has been found, it can then be inserted in all the appropriate places and the data are ready for computing the analysis of variance. The list of treatment totals appears as shown below.

	T
1	$4.77 + x$
2	7.36
3	8.05
4	6.42
5	4.01
6	7.62
7	5.46
8	5.61
9	5.92
	<hr/>
	$G = 55.22 + x$

The quantities needed are

$$T = 4.77, \quad B = 4.02, \quad R = 12.87, \quad G = 55.22$$

$$T_b = 4.77 + 7.36 + 8.05 = 20.18$$

$$B_t = 4.02 + 3.54 + 4.08 + 4.77 = 16.41$$

Hence

$$x = \frac{(9)(4.77) + (12)(4.02) - 12.87 + 55.22 - (3)(20.18) - (3)(16.41)}{12} = 1.98$$

The analysis of variance is now computed in the usual way, except that the total degrees of freedom are reduced from 35 to 34 and the intra-block error degrees of freedom from 16 to 15. The same adjustment factor μ may be used (there is actually a slight change which can be ignored).

When several observations are missing, the values are inserted by the method of successive approximation (illustrated in section 4.25). The exact formulae for t -tests are complicated. The following approximate rule assigns an effective number of replicates to each of two treatments A and B whose means are being compared. In any replicate of the experiment, A is credited with 0 replication if it is absent; with 0 replication if A is present but B occurs in the same block and is absent; and with 1 replication otherwise. The same rule is applied to B . For instance, in a t -test of the difference between treatments (1) and (3) in table 10.1, treatment (1) is credited with 3 replications and treatment (3) with 3 replications (it loses 1 replication because in the first replicate of the experiment treatment (1) is in the same block and is missing). On the other hand, in comparing treatment (1) with treatment (4), the effective numbers of replications are 3 and 4, respectively.

10.2 Partially Balanced Lattices

10.21 Simple Lattices (Two Replicates). For an experiment with 2 replicates, use the first 2 replicates of the appropriate set shown below (plans 10.1–10.6). Notice that designs are available for 36, 100, and 144 treatments (plans 10.7–10.9) as well as for those numbers of treatments for which balanced designs are given. Plans for 121 and 169 treatments can be taken from plans 12.7 and 12.8.

The asymmetry of the designs is apparent from the plans; thus, with 9 treatments (plan 10.1) the first treatment appears in the same block as treatments (2), (3), (4), and (7), but not in the same block as any of the remaining treatments.

For 9 and 16 treatments, simple lattices are unlikely to be more accurate than randomized blocks unless the variation among incomplete blocks is great compared with that within incomplete blocks. Further, the numbers of degrees of freedom for estimating the error are only 4 and 9, as against 9 and 16, respectively, for randomized blocks.

10.22 Triple Lattices (Three Replicates). The first 3 replicates of plans 10.1–10.9, 12.7, and 12.8 are used. Designs may be obtained for all squares from 9 to 169. With 9 treatments the precaution mentioned in section 10.21 applies also to the triple lattice.

10.23 Four Replicates. These designs may be obtained either (i) by duplicating the simple lattice or (ii) by using a quadruple lattice, i.e., the first 4 replicates of the plans. With 36 or 100 treatments, only the first method can be used, since no quadruple lattice exists. The second procedure is slightly preferable, because the resulting design comes closer to symmetry, but the statistical analysis requires more time.

10.24 Five Replicates. The first 5 replicates from the balanced set are taken. Designs are not included for 9, 36, 100, or 144 treatments.

10.25 Higher Numbers of Replicates. Balanced designs should be used for the following numbers of treatments and replicates: 25, 6; 49, 8; 64, 9; and 81, 10. For 9 treatments in 8 or 12 replicates, 16 treatments in 10 replicates, and 25 treatments in 12 replicates, the plan for the balanced design should be repeated.

In other cases the following recommendations are made.

Six replicates. Use the triple lattice *twice*.

Eight replicates. Use the simple lattice *four times* or the quadruple lattice *twice*.

Nine replicates. Use the triple lattice *three times*.

Ten replicates. Use the simple lattice *five times* or the quintuple lattice *twice*.

The designs recommended are not always as fully balanced as they might be made. For instance with 6 replicates of 16 treatments the balanced design (5 replicates) plus 1 extra replicate gives a more nearly symmetrical arrangement than the triple lattice used twice. Since separate computing instructions would be required for such plans, we have preferred to adhere to a more uniform system. For this reason no account is given of designs with 7 replicates, though their statistical analysis presents no great difficulty to the reader who has mastered the principles.

10.26 Arrangement of Experimental Material. In the arrangement of a group of experimental material so as to apply one of those designs, the most important criterion is that units within the same incomplete block be *homogeneous*. In field trials, for example, if the plots are oblong the usual procedure is to have the incomplete blocks as nearly square as practicable, the plots extending the whole length of one side of the block. Uniformity trial investigations have shown that this layout gives on the average the most homogeneous block. In cases where there is detailed knowledge of the experimental site, some other method of grouping may

be superior. Efforts should be made to keep the experimental technique uniform for all units in the same block. Changes in technique that are necessary should be made when changing from one block to another.

When the blocks have been formed, there is a secondary advantage in forming replications so that blocks within the same replicate are as similar to one another as the material permits, since this increases the precision of inter-block comparisons. If the full statistical *analysis is carried out*, it is more important to have the incomplete blocks homogeneous than to have the replications homogeneous. Thus, when the grouping of blocks into compact replications is troublesome, this criterion may be ignored without much loss of precision.

On the other hand, it is desirable to have homogeneous replications if the data may subsequently be analyzed as a randomized block design. As we have pointed out, the randomized block analysis may be needed in fields of work where certain experimental units or whole treatments are likely to be destroyed during the course of the experiment (e.g., in pasture plots subject to winterkill).

10.27 Randomization. The randomization consists of three steps.

1. Randomize the blocks separately and independently within each replication.

2. Randomize the treatments separately and independently within each block.

3. Allot the treatments to the treatment numbers at random.

For methods of randomization, see chapter 15.

Steps 1 and 2 give each treatment an equal chance of being allotted to any experimental unit. These steps correspond to the allotment of treatments to units at random in an ordinary randomized block design.

The function of step 3 is to decide by random choice which groups of treatments will form the blocks of the design. If differences among blocks are large, the error variance per plot for the mean of a group of treatments which lie in the same block may be considerably higher than the average error variance. This additional randomization ensures that the average error variance may be used, in nearly all cases, for comparisons among groups of treatments. For further discussion, see reference (10.4).

When a plan is repeated in order to obtain extra replications, a separate randomization must be made for every replicate.

10.28 Statistical Analysis. An account of the theory, with worked examples of the simple and triple lattice, is given in reference (10.5); a systematic presentation of the shortest computational methods, with worked examples of the simple, quadruple, and balanced lattices in

(10.6); and a detailed description of the methods for carrying out the computations on I.B.M. (Hollerith) punched-card machines in (10.7). The computational methods differ slightly from one reference to another, but all are basically the same.

Table 10.3 shows the plan and yields for a 5×5 simple lattice experiment on soybeans, the treatments being 25 varieties produced in the

TABLE 10.3 YIELDS OF A 5×5 SIMPLE LATTICE EXPERIMENT ON SOYBEANS

(Yields are in bushels per acre, minus 30 bu.)

Rep. I					B	C	μC
(1) 6	(2) 7	(3) 5	(4) 8	(5) 6	32	+ 61	+ 9.5
(6) 16	(7) 12	(8) 12	(9) 13	(10) 8	61	- 8	- 1.3
(11) 17	(12) 7	(13) 7	(14) 9	(15) 14	54	+ 48	+ 7.5
(16) 18	(17) 16	(18) 13	(19) 13	(20) 14	74	- 15	- 2.3
(21) 14	(22) 15	(23) 11	(24) 14	(25) 14	68	+ 17	+ 2.7
					289	+103	+16.1
Rep. II							
(1) 24	(6) 13	(11) 24	(16) 11	(21) 8	80	- 9	- 1.4
(2) 21	(7) 11	(12) 14	(17) 11	(22) 23	80	- 23	- 3.6
(3) 16	(8) 4	(13) 12	(18) 12	(23) 12	56	- 8	- 1.3
(4) 17	(9) 10	(14) 30	(19) 9	(24) 23	89	- 32	- 5.0
(5) 15	(10) 15	(15) 22	(20) 16	(25) 19	87	- 31	- 4.8
					392	-103	-16.1
Treatment totals (unadj.)					μC		
(1) 30	(2) 28	(3) 21	(4) 25	(5) 21	+9.5		
(6) 29	(7) 23	(8) 16	(9) 23	(10) 23	-1.3		
(11) 41	(12) 21	(13) 19	(14) 39	(15) 36	+7.5		
(16) 29	(17) 27	(18) 25	(19) 22	(20) 30	-2.3		
(21) 22	(22) 38	(23) 23	(24) 37	(25) 33	+2.7		
μC -1.4	-3.6	-1.3	-5.0	-4.8	0.0		
Treatment totals (adj.)							
(1) 38.1	(2) 33.9	(3) 29.2	(4) 29.5	(5) 25.7			
(6) 26.3	(7) 18.1	(8) 13.4	(9) 16.7	(10) 16.9			
(11) 47.1	(12) 24.9	(13) 25.2	(14) 41.5	(15) 38.7			
(16) 25.3	(17) 21.1	(18) 21.4	(19) 14.7	(20) 22.9			
(21) 23.3	(22) 37.1	(23) 24.4	(24) 34.7	(25) 30.9			

soybean breeding program of the North Carolina Experiment Station. The experiment actually contained 4 replications, obtained by duplicating the plan for the simple lattice. The first 2 replications will serve to

illustrate the analysis for simple, triple, and quadruple lattices, which all follow the same general pattern. In section 10.29 the data from the whole experiment will be analyzed in order to indicate the procedure when the basic plan is repeated.

1. Find the block totals B , the replication totals, the treatment totals, and the grand total.

2. For each block calculate

$$C = \text{total (over all replicates) of all treatments in the block} - rB$$

For example, for block 1 in replication 1,

$$C = 30 + 28 + 21 + 25 + 21 - (2)(32) = +61$$

Find the replicate totals R_c of the C values. These replicate totals should add to zero.

3. The analysis of variance is as follows.

	d.f.		s.s.	m.s.
Replications	$(r - 1)$	1	212.18	
Treatments (unadj.)	$(k^2 - 1)$	24	559.28	
Blocks within replications (adj.)	$r(k - 1)$	8	501.84	62.73 E_b
Intra-block error	$(k - 1)(rk - k - 1)$	16	218.48	13.66 E_e
Total	$(rk^2 - 1)$	49	1491.78	

All sums of squares are found in the usual way except that for blocks (adjusted), which is given by

$$\frac{\sum C^2}{kr(r - 1)} - \frac{\sum R_c^2}{k^2r(r - 1)} = \frac{(61)^2 + (8)^2 + \cdots + (31)^2}{10} - \frac{(103)^2 + (103)^2}{50} = 501.84$$

where R_c denotes a replication total of the C 's.

4. The weighting factor used to obtain the adjusted treatment totals is

$$\mu = \frac{(E_b - E_e)}{k(r - 1)E_b} = \frac{(62.73 - 13.66)}{(5)(62.73)} = 0.1564$$

where E_b and E_e are, respectively, the mean squares for blocks and intra-block error. If E_b is less than E_e , the factor is taken as zero and no adjustments are made for block effects, the experiment being analyzed as if in randomized blocks.

Each C value is now multiplied by μ to obtain the block corrections μC . These should sum to zero, apart from rounding errors. Each treatment total is adjusted by applying the appropriate correction for every

block in which the treatment appears. For example, the adjusted total for treatment 4 is

$$25 + 9.5 - 5.0 = 29.5$$

In the case of the *simple* lattice it pays to write the μC values around the table of unadjusted treatment totals, putting the values for replication 1 in the right-hand column and those for replication 2 in the bottom row. Then the adjustments to any treatment total are at the end of the row and the bottom of the column in which the treatment lies.

5. The error variance of the difference between two treatment means is slightly smaller for treatments that appear in the same block than for those that do not. The formulae are:

Two treatments in the same block:

$$\frac{2E_e}{r} [1 + (r - 1)\mu] = \frac{(2)(13.66)}{2} [1 + 0.1564] = 15.80$$

Two treatments not in same block:

$$\frac{2E_e}{r} [1 + r\mu] = \frac{(2)(13.66)}{2} [1 + (2)(0.1564)] = 17.93$$

$$\text{Average: } \frac{2E_e}{r} \left[1 + \frac{rk\mu}{(k + 1)} \right] = \frac{(2)(13.66)}{2} \left[1 + \frac{(2)(5)(0.1564)}{6} \right] = 17.22$$

The corresponding standard errors are 3.97, 4.23, and 4.15, respectively. Except with small designs it is sufficient to use the average value, 4.15, for all *t*-tests between pairs of treatments.

6. The analysis of variance does not supply an *F*-test of the adjusted treatment totals. A test of the unadjusted treatment totals can be obtained by analyzing the data as if the experiment were in randomized blocks. The error m.s. is the pooled mean square for blocks and intra-block error, as shown below.

	d.f.		s.s.	m.s.
Treatments (unadj.)	$(k^2 - 1)$	24	559.28	23.30
Error	$(k^2 - 1)(r - 1)$	24	720.32	30.01

This test is not fully sensitive, since it is based on unadjusted treatment totals, but it will be sufficient in cases where an *F*-test is of secondary interest or where differences among incomplete blocks are small.

An *F*-test of the adjusted treatment totals requires a little more calculation. This is most easily done by changing the unadjusted treatments s.s. (which is already available) so that it may be tested against the intra-block error m.s. Calculate B_u , the *unadjusted* sum of squares for blocks within replications, which comes to 350.00. If B_a is the ad-

justed sum of squares for blocks within replications (501.84), we subtract the following quantity from the treatments s.s.

$$k(r-1)\mu \left\{ \left[\frac{r}{(r-1)(1+k\mu)} \right] B_u - B_a \right\} \\ = (5)(1)(0.1564) \left\{ \left[\frac{2}{1+(5)(0.1564)} \right] (350.00) - (501.84) \right\} = -85.30$$

In this case the subtraction term is negative, which is rather unusual. We add 85.30 to the unadjusted treatments s.s. The F -test is then completed by the following analysis.

	d.f.	s.s.	m.s.
Treatments	24	644.58	26.86
Intra-block error	16	218.48	13.66

The F -ratio, 1.97, lies between the 10 and the 5% levels. The F value differs considerably from that obtained by the randomized blocks test, as may happen when the adjustment for incomplete block effects results in a substantial increase in precision.

7. The error m.s. in the randomized blocks analysis was found to be 30.01. To estimate the gain in accuracy over randomized blocks, we compare this figure with the effective error variance

$$E_e \left[1 + \frac{rk\mu}{(k+1)} \right] = (13.66) \left[1 + \frac{(2)(5)(0.1564)}{6} \right] = 17.22$$

The relative accuracy is 30.01/17.22, or 174%.

10.29 Statistical Analysis for Repetitions of the Designs. Suppose that the basic design contains n replicates (where, e.g., $n = 2$ for a simple lattice) and that this is repeated p times, so that the total number of replications is $r = np$. For illustration, we present a joint analysis of the data in tables 10.3 and 10.4 (pp. 275 and 289), which show 4 replications of the 5×5 simple lattice. Replication III is a repetition of replication I, and IV of II. The field results have been rearranged so that treatments follow the same order within corresponding replications. In terms of our notation, $n = 2$, $p = 2$, and $r = 4$.

1. Calculate the block totals, the replication totals, the treatment totals (shown in table 10.4), and the grand total.

2. Arrange the block totals in supplementary tables as on the right of table 10.4. In these tables, blocks that contain the same set of k treatments are placed in the same row. Each table has k rows and p columns, while the number of tables is n . Form the row and column totals of each

table. For each group of similar blocks, compute the quantities

$$C = \text{total (over all replicates) of the treatments appearing in the group} - n(\text{total of the group of blocks})$$

Thus for the first block in replications II and IV, which contains treatments (1), (6), (11), (16), and (21),

$$C = 59 + 51 + 80 + 63 + 58 - 2(157) = -3$$

Find the replication totals R_c of the C values; these should add to zero.

3. The analysis of variance is as follows.

	d.f.		s.s.	m.s.
Replications	$(r - 1)$	3	226.19	
Treatments (unadj.)	$(k^2 - 1)$	24	791.24	
Blocks within replications (adj.)	$r(k - 1)$	16	786.00	$49.12E_b$
Component (a)	$n(p - 1)(k - 1)$	8	164.72	
Component (b)	$n(k - 1)$	8	621.28	
Intra-block error	$(k - 1)(rk - k - 1)$	56	761.56	$13.60E_c$
Total	$(rk^2 - 1)$	99	2564.99	

The blocks s.s. contains two components, both of which are obtained from the supplementary table of block totals.

Component (a) is a new component, which arises only when the design has been repeated. It is composed of the differences between the totals of blocks that contain the same set of treatments. Since each supplementary table has k rows and p columns, it may be analyzed as follows.

	d.f.
Rows	$(k - 1)$
Columns	$(p - 1)$
Rows \times columns	$(k - 1)(p - 1)$
Total	$(kp - 1)$

The sum of squares for component (a) is the sum of the rows \times columns interactions over all n tables. It may be obtained from the following calculations.

$$\text{Total: } \frac{(32)^2 + (61)^2 + \cdots + (75)^2 + (84)^2}{k = 5} - \frac{(650)^2 + (749)^2}{pk^2 = 50} = 602.18$$

$$\text{Rows: } \frac{(104)^2 + (142)^2 + \cdots + (171)^2}{pk = 10} - \frac{(650)^2 + (749)^2}{50} = 309.28$$

$$\text{Columns: } \frac{(289)^2 + (361)^2 + (392)^2 + (357)^2}{k^2 = 25} - \frac{(650)^2 + (749)^2}{50} = 128.18$$

$$\text{Component (a) s.s.} = 602.18 - 309.28 - 128.18 = 164.72$$

The above is a general method that covers all cases. With a simple lattice, a quicker alternative is to calculate the differences between the totals of similar blocks. The sum of squares of deviations of these differences from their replication means, divided by $2k$, gives the sum of squares for component (a).

Component (b) is the component that is present even when there are no repetitions.

$$\begin{aligned}\text{Component (b) s.s.} &= \frac{\sum C^2}{kr(n-1)} - \frac{\sum R_c^2}{k^2r(n-1)} \\ &= \frac{(63)^2 + (29)^2 + \cdots + (48)^2}{20} - \frac{(99)^2 + (99)^2}{100} \\ &= 621.28\end{aligned}$$

The sum of squares for blocks within replications (adjusted) is the pooled sum of squares for the two components. All other terms in the analysis of variance are computed in the usual way.

4. If E_b is the pooled mean square for blocks, and E_e that for error, the weighting factor is

$$\mu = \frac{p(E_b - E_e)}{k[(r-p)E_b + (p-1)E_e]} = \frac{(2)(35.52)}{(5)[(2)(49.12) + 13.60]} = 0.1270$$

where it will be recalled that p is the number of repetitions of the basic design, in this case 2. As usual, no adjustments are made if E_b is less than E_e .

The block corrections μC are obtained and entered in table 10.4; they should add to zero except for rounding errors. Each treatment total is adjusted by applying a correction for each group of blocks in which the treatment appears. For instance, treatment (13) appears in the third block in replications I and III and in the third block in replications II and IV. Its adjusted total is

$$44 + 8.8 - 0.1 = 52.7$$

5. The error variances for the difference between two treatment means are:

Two treatments in the same block:

$$\frac{2E_e}{r}[1 + (n-1)\mu] = \frac{(2)(13.66)}{4}[1 + 0.1270] = 7.70$$

Two treatments not in same block:

$$\frac{2E_e}{r}[1 + n\mu] = \frac{(2)(13.66)}{4}[1 + (2)(0.1270)] = 8.56$$

$$\text{Average: } \frac{2E_e}{r} \left[1 + \frac{nk\mu}{(k+1)} \right] = \frac{(2)(13.66)}{4} \left[1 + \frac{(2)(5)(0.1270)}{6} \right] = 8.28$$

TABLE 10.4 A 5×5 SIMPLE LATTICE (REPS. III AND IV)

Rep. III						Block totals					
						I	III	Sum	C	μC	
(1) 13	(2) 26	(3) 9	(4) 13	(5) 11		72	32	72	104	+63	+ 8.0
(6) 15	(7) 18	(8) 22	(9) 11	(10) 15		81	61	81	142	-29	- 3.7
(11) 19	(12) 10	(13) 10	(14) 10	(15) 16		65	54	65	119	+69	+ 8.8
(16) 21	(17) 16	(18) 17	(19) 4	(20) 17		75	74	75	149	-34	- 4.3
(21) 15	(22) 12	(23) 13	(24) 20	(25) 8		68	68	68	136	+30	+ 3.8
						361	289	361	650	+99	+12.6

Rep. IV						Block totals					
						II	IV	Sum	C	μC	
(1) 16	(6) 7	(11) 20	(16) 13	(21) 21		77	80	77	157	- 3	- 0.4
(2) 15	(7) 10	(12) 11	(17) 7	(22) 14		57	80	57	137	+ 2	+ 0.3
(3) 7	(8) 11	(13) 15	(18) 15	(23) 16		64	56	64	120	- 1	- 0.1
(4) 19	(9) 14	(14) 20	(19) 6	(24) 16		75	89	75	164	-49	- 6.2
(5) 17	(10) 18	(15) 20	(20) 15	(25) 14		84	87	84	171	-48	- 6.1
						357	392	357	749	-99	-12.5

Treatment totals (4 reps.)

(1) 59	(2) 69	(3) 37	(4) 57	(5) 49	+8.0
(6) 51	(7) 51	(8) 49	(9) 48	(10) 56	-3.7
(11) 80	(12) 42	(13) 44	(14) 69	(15) 72	+8.8
(16) 63	(17) 50	(18) 57	(19) 32	(20) 62	-4.3
(21) 58	(22) 64	(23) 52	(24) 73	(25) 55	+3.8
-0.4	+0.3	-0.1	-6.2	-6.1	

Adjusted treatment totals

(1) 66.6	(2) 77.3	(3) 44.9	(4) 58.8	(5) 50.9
(6) 46.9	(7) 47.6	(8) 45.2	(9) 38.1	(10) 46.2
(11) 88.4	(12) 51.1	(13) 52.7	(14) 71.6	(15) 74.7
(16) 58.3	(17) 46.0	(18) 52.6	(19) 21.5	(20) 51.6
(21) 61.4	(22) 68.1	(23) 55.7	(24) 70.6	(25) 52.7

6. An approximate F -test of the adjusted treatment totals is obtained by the method described in the previous section, step 6. In this case the quantity that must be subtracted from the treatments s.s. is

$$k(n-1)\mu \left[\frac{n}{(n-1)(1+k\mu)} B_u - B_a \right]$$

where B_u is the unadjusted and B_a the adjusted sum of squares for component (b) of the blocks. Both quantities have already been computed: B_a , 621.28, appears in the analysis of variance, while B_u , 309.28, is the rows s.s. used in computing component (a).

treatment (14) in replication II. Denote the estimates of the 3 missing values by x , y , and z , respectively; these will be obtained by the method of successive approximation.

It is best to start by finding all block, replicate, and treatment totals and all C values, as in steps 1 and 2 of the computing instructions for this design. Wherever a missing observation is involved, include the appropriate x , y , or z in its place. The summary totals are shown in table 10.5. Note that the total of the C values over the whole experiment is identically zero, so that a good check is available on this part of the calculations.

We must now find first approximations for y and z . In the replications in which they are present, their values are 14 and 9, respectively. However, replication II appears to have higher values than replication I, and it is probably worth while to make an adjustment for replication effect. If we ignore the missing values, the means per plot are about 12 for replication I and 15 for replication II. Since y is missing in replication I, we subtract 3 from its value in replication II, giving 11. For z we add 3, giving 12 as the first approximation. Using $y = 11$, $z = 12$, we now solve for x from formula (10.2). For $r = 2$, $k = 5$ the formula becomes

$$x = \frac{25T - 2R + G - 10C + 5C'}{16}$$

Put $y = 11$, $z = 12$. From table 10.5 we find for x

$$T = 24; \quad R = 276 + 11 = 287; \quad G = 638 + 11 + 12 = 661$$

$$C = 67; \quad C' = 67 - 15 = 52$$

$$x = \frac{(25)(24) - (2)(287) + 661 - (10)(67) + (5)(52)}{16} = \frac{277}{16} = 17$$

Put $x = 17$, $z = 12$. For y we have

$$T = 14; \quad R = 276 + 17 + 293; \quad G = 638 + 17 + 12 = 667$$

$$C = 25 + 12 = 37; \quad C' = 25 + 12 - 30 = 7$$

$$y = \frac{(25)(14) - (2)(293) + 667 - (10)(37) + (5)(7)}{16} = \frac{96}{16} = 6$$

Put $x = 17$, $y = 6$. For z we have

$$T = 9; \quad R = 362; \quad G = 638 + 17 + 6 = 661; \quad C = -2$$

$$C' = -2 + 25 - 6 = 17$$

$$z = \frac{(25)(9) - (2)(362) + 661 - (10)(-2) + (5)(17)}{16} = \frac{267}{16} = 17$$

7. A comparison with randomized blocks is made as in the previous section, step 7. The effective error variance of the lattice is taken as

$$E_e \left[1 + \frac{nk\mu}{(k+1)} \right]$$

10.210 Missing Data. If the missing observations are numerous, an analysis by the method for randomized blocks is recommended. If it is desired to carry out the full analysis, missing values are estimated by the formula which minimizes the intra-block error s.s., though as explained in section 10.13 this method is only approximate.

Experiments with no repetition of the basic design.

$$x = \frac{(r-1)k^2T - rR + G - rkC + kC'}{(r-1)(k-1)(rk-k-1)} \quad (10.2)$$

In this formula T and R are the totals for the treatment and replicate that contain the missing value, G is the grand total, and C is the C value (defined in section 10.28) for the block which contains the missing observation. The only unfamiliar quantity is C' :

C' = total of the C values for all blocks which contain the treatment that has the missing value

Example. For the experiment analyzed in section 10.28, suppose that 3 observations are missing: treatments (1) and (12) in replication I and

TABLE 10.5 SUMMARY TOTALS FOR A 5×5 SIMPLE LATTICE WITH THREE MISSING VALUES

Treatment totals									
(1) 24 + x	(2) 28	(3) 21	(4) 25	(5) 21	(6) 29	(7) 23	(8) 16	(9) 23	(10) 23
(11) 41	(12) 14 + y	(13) 19	(14) 9 + z	(15) 36	(16) 29	(17) 27	(18) 25	(19) 22	(20) 30
(21) 22	(22) 38	(23) 23	(24) 37	(25) 33					

Rep. I

B	C
26 + x	67 - x
61	-8
47 + y	25 - y + z
74	-15
68	17

Rep. II

B	C
80	-15 + x
80	-30 + y
56	-8
59 + z	-2 - z
87	-31

$$R_1 = 276 + x + y \quad 86 - x - y + z \quad R_2 = 362 + z \quad -86 + x + y - z$$

$$G = R_1 + R_2 = 638 + x + y + z$$

treatment (14) in replication II. Denote the estimates of the 3 missing values by x , y , and z , respectively; these will be obtained by the method of successive approximation.

It is best to start by finding all block, replicate, and treatment totals and all C values, as in steps 1 and 2 of the computing instructions for this design. Wherever a missing observation is involved, include the appropriate x , y , or z in its place. The summary totals are shown in table 10.5. Note that the total of the C values over the whole experiment is identically zero, so that a good check is available on this part of the calculations.

We must now find first approximations for y and z . In the replications in which they are present, their values are 14 and 9, respectively. However, replication II appears to have higher values than replication I, and it is probably worth while to make an adjustment for replication effect. If we ignore the missing values, the means per plot are about 12 for replication I and 15 for replication II. Since y is missing in replication I, we subtract 3 from its value in replication II, giving 11. For z we add 3, giving 12 as the first approximation. Using $y = 11$, $z = 12$, we now solve for x from formula (10.2). For $r = 2$, $k = 5$ the formula becomes

$$x = \frac{25T - 2R + G - 10C + 5C'}{16}$$

Put $y = 11$, $z = 12$. From table 10.5 we find for x

$$T = 24; \quad R = 276 + 11 = 287; \quad G = 638 + 11 + 12 = 661$$

$$C = 67; \quad C' = 67 - 15 = 52$$

$$x = \frac{(25)(24) - (2)(287) + 661 - (10)(67) + (5)(52)}{16} = \frac{277}{16} = 17$$

Put $x = 17$, $z = 12$. For y we have

$$T = 14; \quad R = 276 + 17 + 293; \quad G = 638 + 17 + 12 = 667$$

$$C = 25 + 12 = 37; \quad C' = 25 + 12 - 30 = 7$$

$$y = \frac{(25)(14) - (2)(293) + 667 - (10)(37) + (5)(7)}{16} = \frac{96}{16} = 6$$

Put $x = 17$, $y = 6$. For z we have

$$T = 9; \quad R = 362; \quad G = 638 + 17 + 6 = 661; \quad C = -2$$

$$C' = -2 + 25 - 6 = 17$$

$$z = \frac{(25)(9) - (2)(362) + 661 - (10)(-2) + (5)(17)}{16} = \frac{267}{16} = 17$$

This completes the first round. A second round leads to $x = 18$, $y = 5$, $z = 17$, and there is obviously no need for further calculation. The rest of the analysis proceeds as usual except that 3 d.f. are omitted from the total s.s. and the intra-block error s.s. For approximate t -tests, see section 10.13.

Experiments with repetitions of the basic design. If the basic design has n replicates and these are repeated p times to give $r = np$ replications, the formula becomes

$$x = \frac{(n-1)k^2T + (n-1)rR + G - nkC + kC' - n^2R'}{(n-1)(k-1)(rk - k - 1)} \quad (10.3)$$

All symbols have the same meaning as in (10.2) except that C and C' are now derived from the totals of groups of similar blocks, just as in the statistical analysis for this case in section 10.29. The new quantity R' is defined as

R' = total of all replications that are similar to the replication containing the missing value

Note that this total *includes* the replication with the missing value.

Example. In the 5×5 simple lattice with 4 replications, we have $n = 2$, $r = 4$. If treatment (1) is missing in the first replication, the reader may verify that

$$T = 53; \quad R = 283; \quad G = 1393; \quad C = 69; \quad C' = 69 - 9 = 60$$

$$R' = 283 + 361 = 644$$

This gives

$$\begin{aligned} x &= \frac{(25)(53) + (4)(283) + 1393 - (10)(69) + (5)(60) - (4)(644)}{56} \\ &= \frac{844}{56} = 16 \end{aligned}$$

10.3 Rectangular Lattices

10.31 Description. These designs were developed recently by Harshbarger (10.8) for $k(k+1)$ treatments in blocks of k units. They form a useful addition to the square lattices described in previous sections, since the allowable numbers of treatments, 12, 20, 30, 42, 56, 72, etc., fall about midway between the allowable numbers for square lattices. The statistical analysis is quite similar to that for simple and triple lattices, though it takes more time because the block adjustments are not so

simple as with square lattices. The new designs are less symmetrical than the square lattices, in the sense that there is a greater variation in the accuracy with which two treatment means are compared. It will be recalled that only two standard errors are required for t -tests with the simple and triple lattice—one for two treatments that appear in the same block and one for two treatments that do not. The simple rectangular lattice requires four standard errors, while the triple rectangular lattice requires seven. For practical purposes it appears that these can nearly always be reduced to two.

There are several ways of constructing the designs. One (suggested by G. S. Watson) is by means of a latin square with $(k + 1)$ rows and columns, in which every letter in the leading diagonal is different. When writing down the square we attach a number to all letters except those in the leading diagonal, as illustrated below for a 4×4 square.

A	B1	C2	D3
D4	C	B5	A6
B7	A8	D	C9
C10	D11	A12	B

In the first replication we place in a block all numbers that lie in the same row of the latin square, in the second replication all numbers that lie in the same column, and in the third all numbers that have the same latin letter.

Block	Rep. I	Rep. II	Rep. III
1	<u>1 2 3</u>	<u>1 8 11</u>	<u>1 5 7</u>
2	<u>4 5 6</u>	<u>2 5 12</u>	<u>2 9 10</u>
3	<u>7 8 9</u>	<u>3 6 9</u>	<u>3 4 11</u>
4	<u>10 11 12</u>	<u>4 7 10</u>	<u>6 8 12</u>

The important feature of this arrangement is that no two treatments are in the same block more than once. The use of a latin square with different letters down the leading diagonal ensures that in the third replication the three numbers associated with any letter are all in different rows and columns, and hence have not previously occurred together in a block.

Plans 10.10–10.16 were constructed by this method. By using the first 2 replications from any plan we obtain a rectangular lattice in 2 replications, which by means of repetitions can be used for an experiment in 4, 6, 8, etc., replications. By using all 3 replications of the plan we have designs for 3, 6, 9, etc., replications.

Alternatively (as pointed out by S. S. Shrikhande), we may construct the designs from a balanced lattice with $(k + 1)^2$ treatments. If the first replication is omitted, and if all treatments that appear in any one selected block in the first replicate are omitted in subsequent replications, it is easy to verify that we generate a design for $k(k + 1)$ treatments in which every block is of size k and in which no two treatments appear together more than once in the same block. This method gives any number of replicates up to k , though of course the method fails when no balanced lattice exists, as with 36 treatments. The discussion in this section is limited to simple and triple rectangular lattices.

The method of randomizing is the same as for lattices (section 10.27).

10.32 Statistical Analysis. In each plan, the three basic replications are described as the X , Y , and Z replications. Further, if we take any block in one replication and examine another replication, we find that there is one and only one block in the other replication that has no treatments in common with the chosen block. These two blocks will be called *partners*. Since it is important to be able to distinguish partners in the statistical analysis, each block is denoted in the plan by two symbols, e.g., $Y2$, one to mark the replication and one to indicate the partners. All partners carry the same number: thus the partners of $Y2$ are $X2$ and $Z2$. It is advisable to write the block labels in the notebook in which the original results are recorded.

Example. The example is a triple rectangular lattice for 12 treatments in blocks of 3 ($k = 3$). Artificial data were assembled by taking true treatment effects as shown in table 10.6, and adding to them true block effects as given with each block in table 10.7. Thus the observation 7

TABLE 10.6 TREATMENT EFFECTS IN ARTIFICIAL DATA FOR A 3×4
TRIPLE RECTANGULAR LATTICE

Treatment	True effect	True treatment total
1	14	56
2	7	35
3	2	20
4	0	14
5	3	23
6	11	47
7	9	41
8	16	62
9	8	38
10	1	17
11	5	29
12	6	32

for treatment (10) in block X4 of table 10.7 is obtained by adding the true treatment effect, 1, to the true block effect, 6. Since the average of the true block effects comes out to be $56/12$, or $14/3$, the true treatment totals, over the 3 replicates, will be $(3\tau + 14)$, where τ is the true treat-

TABLE 10.7 PLAN AND OBSERVATIONS FOR A 3×4 RECTANGULAR LATTICE
(Treatment numbers are enclosed in parentheses)

Block symbol	True block effect	Rep. X						Totals		Adjustment factor		
								B	C _X			
X4	6	(10)	7	(12)	12	(11)	11	30	-12	-0.6		
X1	2	(2)	9	(3)	4	(1)	16	29	20	+3.4		
X3	7	(7)	16	(9)	15	(8)	23	54	- 5	-1.6		
X2	0	(4)	0	(5)	3	(6)	11	14	30	+5.4		
								127	33			
Rep. Y								B	C _Y			
Y4	9	(3)	11	(6)	20	(9)	17	48	-32	-4.6		
Y2	3	(1)	17	(11)	8	(8)	19	44	10	+1.4		
Y3	3	(12)	9	(2)	10	(5)	6	25	10	+1.4		
Y1	5	(10)	6	(4)	5	(7)	14	25	0	-0.6		
								142	-12			
Rep. Z								B	C _Z			
Z1	4	(8)	20	(6)	15	(12)	10	45	4	+0.2		
Z2	8	(9)	16	(10)	9	(2)	15	40	-16	-3.8		
Z3	1	(11)	6	(3)	3	(4)	1	10	19	+3.2		
Z4	8	(5)	11	(1)	22	(7)	17	50	-28	-3.8		
								145	-21			
Treatment totals												
	1	2	3	4	5	6	7	8	9	10	11	12
Unadj.	55	34	18	6	20	46	47	62	48	22	25	31
Adj.	56	35	20	14	23	47	41	62	38	17	29	32

ment effect. These figures are shown in the column at the right in table 10.6.

No intra-block error has been introduced into the data, which therefore provide two checks on the method of analysis. First, in the analysis of variance the intra-block error s.s. should be found to be identically zero. Second, the adjusted treatment totals as found from the analysis should be *exactly* equal to the true treatment totals.

The computing instructions given below apply to either the simple or the triple rectangular lattice without repetitions. The changes required when there are repetitions are given later.

1. Find the block totals, B , the replication totals, the treatment totals, and the grand total.

2. For each block calculate

$$C = \text{total (over all replicates) of all treatments in the block} - rB$$

Thus for block Y3,

$$C = 18 + 46 + 48 - (3)(48) = -32$$

As usual, the total of the C 's is zero.

3. Arrange the C values in a supplementary table so that partners appear in the same row. The row totals of this table, S , give the sums of the C values for each set of partners. The column totals give values denoted by R_C .

TABLE 10.8 SUPPLEMENTARY TABLE OF C VALUES

Block symbol	C_X	C_Y	C_Z	S	λC_X	λC_Y	λC_Z	μS
1	20	0	4	24	4.0	0.0	0.8	0.6
2	30	10	-16	24	6.0	2.0	-3.2	0.6
3	-5	10	19	24	-1.0	2.0	3.8	0.6
4	-12	-32	-28	-72	-2.4	-6.4	-5.6	-1.8
Totals (R_C)	33	-12	-21	0	6.6	-2.4	-4.2	0.0

4. The analysis of variance is now obtained. All sums of squares are found in the usual way, except that for blocks, adjusted for treatments, which is

$$\begin{aligned}
 & \frac{\sum C^2}{r(rk - k - 1)} - \frac{\sum R_C^2}{r(k + 1)(rk - k - 1)} - \frac{\sum S^2}{r(r - 1)(k + 1)(rk - k - 1)} \\
 &= \frac{(20)^2 + (30)^2 + \dots + (28)^2}{15} - \frac{(33)^2 + (12)^2 + (21)^2}{60} - \frac{(24)^2 + \dots + (72)^2}{120} \\
 &= 274.0 - 27.9 - 57.6 = 188.5 \quad (10.4)
 \end{aligned}$$

TABLE 10.9 ANALYSIS OF VARIANCE

Source of variation	d.f.	s.s.	m.s.
Replications	$(r - 1)$	2	15.5
Treatments	$(k^2 + k - 1)$	11	1067.0
Blocks	rk	9	188.5
Intra-block error	$(r - 1)(k^2 - 1) - k$	13	0.0
Total	$(rk^2 + rk - 1)$	35	1271.0

As anticipated, the intra-block error s.s. in table 10.9 is zero.

5. We now calculate the weighting factors used to obtain the adjusted treatment totals. Although a general formula can be given, it is simpler to present these separately for simple and triple designs. Two weighting factors are necessary.*

Simple rectangular lattice

$$\lambda = \frac{r(E_b - E_e)}{r(k-1)E_b + (rk - 2k + r)E_e} \quad (10.5)$$

$$\mu = \frac{\lambda r(E_b - E_e)}{r(k+1)E_b + (rk - 2k - r)E_e} \quad (10.6)$$

Triple rectangular lattice

$$\lambda = \frac{r(E_b - E_e)}{r(2k-1)E_b + (rk - 6k + r)E_e} \quad (10.7)$$

$$\mu = \frac{\lambda r(E_b - E_e)}{2r(k+1)E_b + (rk - 6k - 2r)E_e} \quad (10.8)$$

In this example E_e is zero and we have

$$\lambda = \frac{3}{(3)(5)} = 0.2; \quad \mu = \frac{(0.2)(3)}{(2)(3)(4)} = 0.025$$

6. Complete table 10.8 by adding the columns λC_X , λC_Y , λC_Z , and μS . The adjustment for any block is

$$\lambda C - \mu S$$

where S is taken from the row in which C lies in table 10.8. Thus for block $X1$ the adjustment is

$$+4.0 - 0.6 = +3.4$$

These adjustments are recorded on the plan in table 10.7. Note that the order in which blocks appear is different in tables 10.7 and 10.8; care must be taken to see that each adjustment is given to the appropriate block.

7. Finally, each treatment *total* is adjusted by adding the adjustments for every block in which the treatment appears. For instance, the adjusted total for treatment (4) is

$$6 + 5.4 - 0.6 + 3.2 = 14$$

* This method of estimating the weights is slightly different from that given by Harshbarger. In the interests of simplicity we have used only the pooled mean square for blocks and the intra-block error m.s. for estimating λ and μ .

Every adjusted treatment total will be found to be equal to the true treatment total as given in table 10.6.

Statistical analysis for repetitions of the designs. When the basic design with n replications is repeated p times to give $r = np$ replications, the changes required in the steps of the analysis are noted below.

1. No change.

2a. For any block there are $(p - 1)$ other blocks that have exactly the same set of treatments. Arrange the totals for such similar blocks in two-way tables, as exemplified below. There are n such tables.

Block no.	X Blocks Repetition				Total ($\sum \beta$)
	1	2	...	p	
1					
2					
.					
.					
($k + 1$)					
Total					

2b. For each group of similar blocks calculate

$C = \text{total (over all replicates) of all treatments in the group} - n(\sum \beta)$

3. No change.

4a. The blocks s.s. now has two components. For component (a) obtain the interaction s.s. for each of the tables in step 2a, and add these sums of squares. Since each block total contains k observations, a divisor k is required for the analysis of variance.

4b. The sum of squares for component b is

$$\frac{\sum C^2}{r(nk - k - 1)} - \frac{\sum R_c^2}{r(k + 1)(nk - k - 1)} - \frac{\sum S^2}{r(n - 1)(k + 1)(nk - k - 1)}$$

The separation of degrees of freedom in the analysis of variance is shown below.

	d.f.	m.s.
Replications	$(r - 1)$	
Treatments	$(k^2 + k - 1)$	
Blocks	rk	E_b
Component (a)	$(r - n)k$	
Component (b)	nk	
Intra-block error	$(r - 1)(k^2 - 1) - k$	E_e
Total	$(rk^2 + rk - 1)$	

5. The weighting factors in formulae (10.5)–(10.8) are unchanged. Note that E_b is the pooled mean square for blocks.

6. Unchanged.

7. Unchanged.

As mentioned previously, in order to have t -tests for every pair of treatments we would require to present 4 standard error formulae for simple rectangular lattices and 7 for triple rectangular lattices. With very little inaccuracy these can be reduced to 2, one for the case where the two treatments appear in a block and one for the case where this is not so. For the difference between two adjusted *means* the estimated error variances are:

Simple rectangular lattice

Two treatments in the same block: $\frac{2E_e}{r}(1 + \lambda - \mu)$

Two treatments not in same block: $\frac{2E_e}{r}(1 + 2\lambda - 3\mu)$

$$\begin{aligned} \text{Average: } * \frac{2E_e}{r} \left[\frac{2(k-1)}{k^2+k-1}(1 + \lambda - \mu) + \frac{k^2-k+1}{k^2+k-1}(1 + 2\lambda - 3\mu) \right] \\ = \frac{2E_e}{r} \left[1 + \frac{2k^2\lambda - (k^2+k-1)\mu}{(k^2+k-1)} \right] \end{aligned}$$

Triple rectangular lattice

Two treatments in the same block: $\frac{2E_e}{r}(1 + 2\lambda - \mu)$

Two treatments not in same block: $\frac{2E_e}{r} \left(1 + 3\lambda - \frac{3}{2}\mu \right)$

$$\text{Average: } * \frac{2E_e}{r} \left[\frac{3(k-1)}{k^2+k-1}(1 + 2\lambda - \mu) + \frac{k^2-2k+2}{k^2+k-1} \left(1 + 3\lambda - \frac{3}{2}\mu \right) \right]$$

In the example, where $\lambda = 0.2$, $\mu = 0.025$, the factors multiplying the usual term $2E_e/r$ are 1.375 for two treatments in the same block and 1.562 for two treatments not in the same block. The factor in the average variance is 1.460.

10.4 Cubic Lattices

10.41 Description. These designs were produced by Yates for plant-breeding work in which selections are to be made from an unusually large number of varieties. The number of treatments must be an exact cube.

* Owing to the fact that only two standard errors were used, this average is not quite equal to the average variance taken over all possible pairs, though it is very close to that value.

The most useful range comprizes 27, 64, 125, 216, 343, 512, 729, and 1000 treatments. The size of block is the cube root of the number of treatments, i.e., 3, 4, 5, 6, 7, 8, 9, and 10, respectively. Thus cubic lattices can accommodate a large number of treatments in a small size of incomplete block. The designs have been used, for example, in an experiment with 729 strains of *ponderosa* pine seedlings (10.9) and an experiment with 729 soybean varieties (10.10). The number of replicates must be 3 or some multiple of 3.

Since the plans are easy to construct and since they occupy a considerable amount of space for the higher numbers of treatments, they are not reproduced here. To obtain a plan, the k^3 treatments are numbered by means of a three-digit code in which each digit takes all values from 1 to k . For 27 treatments, the code is as follows:

T^*	Code	T'	Code	T''	Code
1	111	4	121	7	131
2	211	5	221	8	231
3	311	6	321	9	331
10	112	13	122	16	132
11	212	14	222	17	232
12	312	15	322	18	332
19	113	22	123	25	133
20	213	23	223	26	233
21	313	24	323	27	333

* Treatment number.

The same principle applies with a larger number of treatments. For the first k treatments, the last two digits are fixed at (11) while the first digit runs from 1 to k . The next k treatments are coded by fixing the last two digits at (21) while the first digit again runs from 1 to k , and so on in a systematic manner until the final k treatments are reached, for which the last two digits have the fixed values (kk).

Within each of the 3 replications, the k^3 treatments are grouped into k^2 blocks, each of size k . In the first replication, the rule for this grouping is to keep the last two digits constant within a block, allowing the first digit to take all values from 1 to k . Thus, in the example above, the 9 groups of treatments constitute the 9 blocks, block 1 containing the treatments (111), (211), and (311).

To form blocks in the second replication, we keep the first and last digits fixed within any block and give the second digit all values from 1 to k . With 27 treatments, the first block therefore contains (111), (121), and (131), the second block (211), (221), and (231), and the last

block (313), (323), and (333). In the third replication, the first and second digits are constant within each block.

The composition of the blocks in the second and third replications is shown in table 10.10.

TABLE 10.10 SECOND AND THIRD REPLICATES OF A CUBIC LATTICE WITH 27 TREATMENTS

Rep. II										
Block	(1)		(2)		(3)		(4)		(5)	
	Code	<i>T</i> *	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>
	111	1	211	2	311	3	112	10	212	11
	121	4	221	5	321	6	122	13	222	14
	131	7	231	8	331	9	132	16	232	17
	(6)		(7)		(8)		(9)			
	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>		
	312	12	113	19	213	20	313	21		
	322	15	123	22	223	23	323	24		
	332	18	133	25	233	26	333	27		
Rep. III										
Block	(1)		(2)		(3)		(4)		(5)	
	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>
	111	1	211	2	311	3	121	4	221	5
	112	10	212	11	312	12	122	13	222	14
	113	19	213	20	313	21	123	22	223	23
	(6)		(7)		(8)		(9)			
	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>	Code	<i>T</i>		
	321	6	131	7	231	8	331	9		
	322	15	132	16	232	17	332	18		
	323	24	133	25	233	26	333	27		

* Treatment number.

The original treatment numbers are shown as well as the code numbers. In an experiment it is simplest to record only the code numbers, which are required in order to follow the computing instructions.

Cubic lattices may be expected to be most useful when the number of treatments exceeds 100. The designs for $k = 3, 4, 5$, and 6 lie in the range which is also covered by lattice designs. We do not know of any investigations of the relative accuracy of lattice and cubic lattice designs in these cases. With highly variable experimental material, the use of a smaller block might give the cubic lattice some advantage; on the other hand the statistical analysis is more laborious owing to the greater num-

ber of block adjustments to be calculated. Like the lattice designs, cubic lattices cannot be appreciably less accurate than randomized blocks which occupy the same set of replications, and can be analyzed by the method for randomized blocks.

Some cubic lattices can also be arranged in quasi-latin squares (10.11) so as to allow the elimination of two sources of error variation.

10.42 Arrangement of Experimental Material. The most important rule is to have units in the same incomplete block as homogeneous as the experimental material permits; in field trials the blocks should be compact in shape. If practicable, blocks within the same replicate should also be similar to each other, so as to increase the accuracy of inter-block comparisons and to obtain more precise results if the analysis is subsequently carried out by the method for randomized blocks.

10.43 Randomization. The steps are:

1. Randomize the order of the blocks independently within each replication.
2. Randomize the positions of the treatment code numbers independently within each block.
3. Assign treatments to code numbers at random.

10.44 Statistical Analysis. The following reference should be consulted.

- (10.12) YATES, F. The recovery of inter-block information in variety trials arranged in three-dimensional lattices. *Ann. Eugen.* 9, 136-156, 1939.

General formulae are given with a numerical example of the analysis for an experiment with 64 treatments. In following this example, the reader should note that the experiment was not arranged in separate replications; the changes introduced for this reason are pointed out by Yates (pp. 147-148). The method of analysis when 6 or 9 replicates are used is also indicated (p. 144). Reference (10.13) describes the analysis of an experiment with 125 treatments.

10.45 Error Variances. The formulae for the error variance of the difference between two treatment *means* are summarized below. In the analysis as described by Yates (10.12), the symbols

$$\lambda = \frac{w - w'}{w + 2w'}, \quad \mu = \frac{w - w'}{2w + w'}$$

play a prominent part. It is convenient to write $\lambda' = \lambda/k^2$, $\mu' = \mu/k^2$. The error variances for the different types of treatment comparisons are as follows:

Comparison	Variance of difference between means
$t_{211} - t_{111}$	$\frac{2E_e}{r} [1 + 2\lambda' + 2(k-1)\mu']$
$t_{122} - t_{111}$	$\frac{2E_e}{r} [1 + 4\lambda' + (3k-4)\mu']$
$t_{222} - t_{111}$	$\frac{2E_e}{r} [1 + 6\lambda' + 3(k-2)\mu']$
Average	$\frac{2E_e}{r} \left[1 + \frac{k^2}{(k^2 + k + 1)} \{6\lambda' + 3(k-1)\mu'\} \right]$

Here E_e is the intra-block error m.s. and r the number of replicates (3, 6, or 9). The formulae, though written in a slightly different form, are identical with those given by Yates. Our k corresponds to Yates's p , and our r to his n .

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PLANS

Plan 10.1

 3×3 balanced lattice

$$t = 9, k = 3, r = 4, b = 12, \lambda = 1^*$$

Block	Rep. I	Rep. II	Rep. III	Rep. IV
(1)	<u>1 2 3</u>	(4) <u>1 4 7</u>	(7) <u>1 5 9</u>	(10) <u>1 8 6</u>
(2)	<u>4 5 6</u>	(5) <u>2 5 8</u>	(8) <u>7 2 6</u>	(11) <u>4 2 9</u>
(3)	<u>7 8 9</u>	(6) <u>3 6 9</u>	(9) <u>4 8 3</u>	(12) <u>7 5 3</u>

Plan 10.2

 4×4 balanced lattice

$$t = 16, k = 4, r = 5, b = 20, \lambda = 1$$

Block	Rep. I	Rep. II	Rep. III
(1)	<u>1 2 3 4</u>	(5) <u>1 5 9 13</u>	(9) <u>1 6 11 16</u>
(2)	<u>5 6 7 8</u>	(6) <u>2 6 10 14</u>	(10) <u>5 2 15 12</u>
(3)	<u>9 10 11 12</u>	(7) <u>3 7 11 15</u>	(11) <u>9 14 3 8</u>
(4)	<u>13 14 15 16</u>	(8) <u>4 8 12 16</u>	(12) <u>13 10 7 4</u>

	Rep. IV	Rep. V
(13)	<u>1 14 7 12</u>	(17) <u>1 10 15 8</u>
(14)	<u>13 2 11 8</u>	(18) <u>9 2 7 16</u>
(15)	<u>5 10 3 16</u>	(19) <u>13 6 3 12</u>
(16)	<u>9 6 15 4</u>	(20) <u>5 14 11 4</u>

Plan 10.3

 5×5 balanced lattice

$$t = 25, k = 5, r = 6, b = 30, \lambda = 1$$

Block	Rep. I	Rep. II	Rep. III
(1)	<u>1 2 3 4 5</u>	(6) <u>1 6 11 16 21</u>	(11) <u>1 7 13 19 25</u>
(2)	<u>6 7 8 9 10</u>	(7) <u>2 7 12 17 22</u>	(12) <u>21 2 8 14 20</u>
(3)	<u>11 12 13 14 15</u>	(8) <u>3 8 13 18 23</u>	(13) <u>16 22 3 9 15</u>
(4)	<u>16 17 18 19 20</u>	(9) <u>4 9 14 19 24</u>	(14) <u>11 17 23 4 10</u>
(5)	<u>21 22 23 24 25</u>	(10) <u>5 10 15 20 25</u>	(15) <u>6 12 18 24 5</u>

	Rep. IV	Rep. V	Rep. VI
(16)	<u>1 12 23 9 20</u>	(21) <u>1 17 8 24 15</u>	(26) <u>1 22 18 14 10</u>
(17)	<u>16 2 13 24 10</u>	(22) <u>11 2 18 9 25</u>	(27) <u>6 2 23 19 15</u>
(18)	<u>6 17 3 14 25</u>	(23) <u>21 12 3 19 10</u>	(28) <u>11 7 3 24 20</u>
(19)	<u>21 7 18 4 15</u>	(24) <u>6 22 13 4 20</u>	(29) <u>16 12 8 4 25</u>
(20)	<u>11 22 8 19 5</u>	(25) <u>16 7 23 14 5</u>	(30) <u>21 17 13 9 5</u>

*The symbol λ denotes the number of times that two treatments appear in the same block.

Plan 10.4

 7×7 balanced lattice

$$t = 49, k = 7, r = 8, b = 56, \lambda = 1$$

Block	Rep. I							Rep. II							
(1)	1	2	3	4	5	6	7	(8)	1	8	15	22	29	36	43
(2)	8	9	10	11	12	13	14	(9)	2	9	16	23	30	37	44
(3)	15	16	17	18	19	20	21	(10)	3	10	17	24	31	38	45
(4)	22	23	24	25	26	27	28	(11)	4	11	18	25	32	39	46
(5)	29	30	31	32	33	34	35	(12)	5	12	19	26	33	40	47
(6)	36	37	38	39	40	41	42	(13)	6	13	20	27	34	41	48
(7)	43	44	45	46	47	48	49	(14)	7	14	21	28	35	42	49
Rep. III							Rep. IV								
(15)	1	9	17	25	33	41	49	(22)	1	37	24	11	47	34	21
(16)	43	2	10	18	26	34	42	(23)	15	2	38	25	12	48	35
(17)	36	44	3	11	19	27	35	(24)	29	16	3	39	26	13	49
(18)	29	37	45	4	12	20	28	(25)	43	30	17	4	40	27	14
(19)	22	30	38	46	5	13	21	(26)	8	44	31	18	5	41	28
(20)	15	23	31	39	47	6	14	(27)	22	9	45	32	19	6	42
(21)	8	16	24	32	40	48	7	(28)	36	23	10	46	33	20	7
Rep. V							Rep. VI								
(29)	1	30	10	39	19	48	28	(36)	1	23	45	18	40	13	35
(30)	22	2	31	11	40	20	49	(37)	29	2	24	46	19	41	14
(31)	43	23	3	32	12	41	21	(38)	8	30	3	25	47	20	42
(32)	15	44	24	4	33	13	42	(39)	36	9	31	4	26	48	21
(33)	36	16	45	25	5	34	14	(40)	15	37	10	32	5	27	49
(34)	8	37	17	46	26	6	35	(41)	43	16	38	11	33	6	28
(35)	29	9	38	18	47	27	7	(42)	22	44	17	39	12	34	7
Rep. VII							Rep. VIII								
(43)	1	16	31	46	12	27	42	(50)	1	44	38	32	26	20	14
(44)	36	2	17	32	47	13	28	(51)	8	2	45	39	33	27	21
(45)	22	37	3	18	33	48	14	(52)	15	9	3	46	40	34	28
(46)	8	23	38	4	19	34	49	(53)	22	16	10	4	47	41	35
(47)	43	9	24	39	5	20	35	(54)	29	23	17	11	5	48	42
(48)	29	44	10	25	40	6	21	(55)	36	30	24	18	12	6	49
(49)	15	30	45	11	26	41	7	(56)	43	37	31	25	19	13	7

Plan 10.5

 8×8 balanced lattice

$$t = 64, k = 8, r = 9, b = 72, \lambda = 1$$

Block	Rep. I							
(1)	1	2	3	4	5	6	7	8
(2)	9	10	11	12	13	14	15	16
(3)	17	18	19	20	21	22	23	24
(4)	25	26	27	28	29	30	31	32
(5)	33	34	35	36	37	38	39	40
(6)	41	42	43	44	45	46	47	48
(7)	49	50	51	52	53	54	55	56
(8)	57	58	59	60	61	62	63	64

	Rep. III							
(17)	1	10	19	28	37	46	55	64
(18)	9	2	51	44	61	30	23	40
(19)	17	50	3	36	29	62	15	48
(20)	25	42	35	4	21	14	63	56
(21)	33	58	27	20	5	54	47	16
(22)	41	26	59	12	53	6	39	24
(23)	49	18	11	60	45	38	7	32
(24)	57	34	43	52	13	22	31	8

	Rep. V							
(33)	1	26	43	60	21	54	15	40
(34)	25	2	11	52	37	62	47	24
(35)	41	10	3	20	61	38	31	56
(36)	57	50	19	4	45	30	39	16
(37)	17	34	59	44	5	14	55	32
(38)	49	58	35	28	13	6	23	48
(39)	9	42	27	36	53	22	7	64
(40)	33	18	51	12	29	46	63	8

	Rep. II							
(9)	1	9	17	25	33	41	49	57
(10)	2	10	18	26	34	42	50	58
(11)	3	11	19	27	35	43	51	59
(12)	4	12	20	28	36	44	52	60
(13)	5	13	21	29	37	45	53	61
(14)	6	14	22	30	38	46	54	62
(15)	7	15	23	31	39	47	55	63
(16)	8	16	24	32	40	48	56	64

	Rep. IV							
(25)	1	18	27	44	13	62	39	56
(26)	17	2	35	60	53	46	31	16
(27)	25	34	3	12	45	54	23	64
(28)	41	58	11	4	29	22	55	40
(29)	9	50	43	28	5	38	63	24
(30)	57	42	51	20	37	6	15	32
(31)	33	26	19	52	61	14	7	48
(32)	49	10	59	36	21	30	47	8

	Rep. VI							
(41)	1	34	11	20	53	30	63	48
(42)	33	2	59	28	45	22	15	56
(43)	9	58	3	52	21	46	39	32
(44)	17	26	51	4	13	38	47	64
(45)	49	42	19	12	5	62	31	40
(46)	25	18	43	36	61	6	55	16
(47)	57	10	35	44	29	54	7	24
(48)	41	50	27	60	37	14	23	8

Plan 10.5 (Continued)

8 × 8 balanced lattice

Rep. VII							
(49)	1	42	59	52	29	38	23 16
(50)	41	2	19	36	13	54	63 32
(51)	57	18	3	28	53	14	47 40
(52)	49	34	27	4	61	46	15 24
(53)	25	10	51	60	5	22	39 48
(54)	33	50	11	44	21	6	31 64
(55)	17	58	43	12	37	30	7 56
(56)	9	26	35	20	45	62	55 8

Rep. IX							
(65)	1	58	51	36	45	14	31 24
(66)	57	2	27	12	21	38	55 48
(67)	49	26	3	44	37	22	63 16
(68)	33	10	43	4	53	62	23 32
(69)	41	18	35	52	5	30	15 64
(70)	9	34	19	60	29	6	47 56
(71)	25	50	59	20	13	46	7 40
(72)	17	42	11	28	61	54	39 8

Rep. VIII							
(57)	1	50	35	12	61	22	47 32
(58)	49	2	43	20	29	14	39 64
(59)	33	42	3	60	13	30	55 24
(60)	9	18	59	4	37	54	31 48
(61)	57	26	11	36	5	46	23 56
(62)	17	10	27	52	45	6	63 40
(63)	41	34	51	28	21	62	7 16
(64)	25	58	19	44	53	38	15 8

Plan 10.6

9 × 9 balanced lattice

$$t = 81, k = 9, r = 10, b = 90, \lambda = 1$$

Block	Rep. I								
(1)	1	2	3	4	5	6	7	8	9
(2)	10	11	12	13	14	15	16	17	18
(3)	19	20	21	22	23	24	25	26	27
(4)	28	29	30	31	32	33	34	35	36
(5)	37	38	39	40	41	42	43	44	45
(6)	46	47	48	49	50	51	52	53	54
(7)	55	56	57	58	59	60	61	62	63
(8)	64	65	66	67	68	69	70	71	72
(9)	73	74	75	76	77	78	79	80	81

Rep. III								
(19)	1	20	12	58	77	69	34	53 45
(20)	10	2	21	67	59	78	43	35 54
(21)	19	11	3	76	68	60	52	44 36
(22)	28	47	39	4	23	15	61	80 72
(23)	37	29	48	13	5	24	70	62 81
(24)	46	38	30	22	14	6	79	71 63
(25)	55	74	66	31	50	42	7	26 18
(26)	64	56	75	40	32	51	16	8 27
(27)	73	65	57	49	41	33	25	17 9

Rep. II								
(10)	1	10	19	28	37	46	55	64 73
(11)	2	11	20	29	38	47	56	65 74
(12)	3	12	21	30	39	48	57	66 75
(13)	4	13	22	31	40	49	58	67 76
(14)	5	14	23	32	41	50	59	68 77
(15)	6	15	24	33	42	51	60	69 78
(16)	7	16	25	34	43	52	61	70 79
(17)	8	17	26	35	44	53	62	71 80
(18)	9	18	27	36	45	54	63	72 81

Rep. IV								
(28)	1	11	21	31	41	51	61	71 81
(29)	19	2	12	49	32	42	79	62 72
(30)	10	20	3	40	50	33	70	80 63
(31)	55	65	75	4	14	24	34	44 54
(32)	73	56	66	22	5	15	52	35 45
(33)	64	74	57	13	23	6	43	53 36
(34)	28	38	48	58	68	78	7	17 27
(35)	46	29	39	76	59	69	25	8 18
(36)	37	47	30	67	77	60	16	26 9

Plan 10.6 (Continued)

9 × 9 balanced lattice

Rep. V

(37)	1	29	57	22	50	78	16	44	72
(38)	55	2	30	76	23	51	70	17	45
(39)	28	56	3	49	77	24	43	71	18
(40)	10	38	66	4	32	60	25	53	81
(41)	64	11	39	58	5	33	79	26	54
(42)	37	65	12	31	59	6	52	80	27
(43)	19	47	75	13	41	69	7	35	63
(44)	73	20	48	67	14	42	61	8	36
(45)	46	74	21	40	68	15	34	62	9

Rep. VI

(46)	1	56	30	13	68	42	25	80	54
(47)	28	2	57	40	14	69	52	26	81
(48)	55	29	3	67	41	15	79	53	27
(49)	19	74	48	4	59	33	16	71	45
(50)	46	20	75	31	5	60	43	17	72
(51)	73	47	21	58	32	6	70	44	18
(52)	10	65	39	22	77	51	7	62	36
(53)	37	11	66	49	23	78	34	8	63
(54)	64	38	12	76	50	24	61	35	9

Rep. VII

(55)	1	47	66	76	14	33	43	62	27
(56)	64	2	48	31	77	15	25	44	63
(57)	46	65	3	13	32	78	61	26	45
(58)	37	56	21	4	50	69	79	17	36
(59)	19	38	57	67	5	51	34	80	18
(60)	55	20	39	49	68	6	16	35	81
(61)	73	11	30	40	59	24	7	53	72
(62)	28	74	12	22	41	60	70	8	54
(63)	10	29	75	58	23	42	52	71	9

Rep. VIII

(64)	1	74	39	67	32	24	52	17	63
(65)	37	2	75	22	68	33	61	53	18
(66)	73	38	3	31	23	69	16	62	54
(67)	46	11	57	4	77	42	70	35	27
(68)	55	47	12	40	5	78	25	71	36
(69)	10	56	48	76	41	6	34	26	72
(70)	64	29	21	49	14	60	7	80	45
(71)	19	65	30	58	50	15	43	8	81
(72)	28	20	66	13	59	51	79	44	9

Rep. IX

(73)	1	65	48	40	23	60	79	35	18
(74)	46	2	66	58	41	24	16	80	36
(75)	64	47	3	22	59	42	34	17	81
(76)	73	29	12	4	68	51	43	26	63
(77)	10	74	30	49	5	69	61	44	27
(78)	28	11	75	67	50	6	25	62	45
(79)	37	20	57	76	32	15	7	71	54
(80)	55	38	21	13	77	33	52	8	72
(81)	19	56	39	31	14	78	70	53	9

Rep. X

(82)	1	38	75	49	59	15	70	26	36
(83)	73	2	39	13	50	60	34	71	27
(84)	37	74	3	58	14	51	25	35	72
(85)	64	20	30	4	41	78	52	62	18
(86)	28	65	21	76	5	42	16	53	63
(87)	19	29	66	40	77	6	61	17	54
(88)	46	56	12	67	23	33	7	44	81
(89)	10	47	57	31	68	24	79	8	45
(90)	55	11	48	22	32	69	43	80	9

Plan 10.7

 6×6 triple lattice

Block	Rep. I						Rep. II						
(1)	1	2	3	4	5	6	(7)	1	7	13	19	25	31
(2)	7	8	9	10	11	12	(8)	2	8	14	20	26	32
(3)	13	14	15	16	17	18	(9)	3	9	15	21	27	33
(4)	19	20	21	22	23	24	(10)	4	10	16	22	28	34
(5)	25	26	27	28	29	30	(11)	5	11	17	23	29	35
(6)	31	32	33	34	35	36	(12)	6	12	18	24	30	36

Rep. III						
(13)	1	8	15	22	29	36
(14)	31	2	9	16	23	30
(15)	25	32	3	10	17	24
(16)	19	26	33	4	11	18
(17)	13	20	27	34	5	12
(18)	7	14	21	28	35	6

Plan 10.8

 10×10 triple lattice

Block	Rep. I										Rep. II										
(1)	1	2	3	4	5	6	7	8	9	10	(11)	1	11	21	31	41	51	61	71	81	91
(2)	11	12	13	14	15	16	17	18	19	20	(12)	2	12	22	32	42	52	62	72	82	92
(3)	21	22	23	24	25	26	27	28	29	30	(13)	3	13	23	33	43	53	63	73	83	93
(4)	31	32	33	34	35	36	37	38	39	40	(14)	4	14	24	34	44	54	64	74	84	94
(5)	41	42	43	44	45	46	47	48	49	50	(15)	5	15	25	35	45	55	65	75	85	95
(6)	51	52	53	54	55	56	57	58	59	60	(16)	6	16	26	36	46	56	66	76	86	96
(7)	61	62	63	64	65	66	67	68	69	70	(17)	7	17	27	37	47	57	67	77	87	97
(8)	71	72	73	74	75	76	77	78	79	80	(18)	8	18	28	38	48	58	68	78	88	98
(9)	81	82	83	84	85	86	87	88	89	90	(19)	9	19	29	39	49	59	69	79	89	99
(10)	91	92	93	94	95	96	97	98	99	100	(20)	10	20	30	40	50	60	70	80	90	100

Rep. III										
(21)	1	12	23	34	45	56	67	78	89	100
(22)	91	2	13	24	35	46	57	68	79	90
(23)	81	92	3	14	25	36	47	58	69	80
(24)	71	82	93	4	15	26	37	48	59	70
(25)	61	72	83	94	5	16	27	38	49	60
(26)	51	62	73	84	95	6	17	28	39	50
(27)	41	52	63	74	85	96	7	18	29	40
(28)	31	42	53	64	75	86	97	8	19	30
(29)	21	32	43	54	65	76	87	98	9	20
(30)	11	22	33	44	55	66	77	88	99	10

Plan 10.9

12 × 12 quadruple lattice

Block	Rep. I											
(1)	1	2	3	4	5	6	7	8	9	10	11	12
(2)	13	14	15	16	17	18	19	20	21	22	23	24
(3)	25	26	27	28	29	30	31	32	33	34	35	36
(4)	37	38	39	40	41	42	43	44	45	46	47	48
(5)	49	50	51	52	53	54	55	56	57	58	59	60
(6)	61	62	63	64	65	66	67	68	69	70	71	72
(7)	73	74	75	76	77	78	79	80	81	82	83	84
(8)	85	86	87	88	89	90	91	92	93	94	95	96
(9)	97	98	99	100	101	102	103	104	105	106	107	108
(10)	109	110	111	112	113	114	115	116	117	118	119	120
(11)	121	122	123	124	125	126	127	128	129	130	131	132
(12)	133	134	135	136	137	138	139	140	141	142	143	144

	Rep. II											
(13)	1	13	25	37	49	61	73	85	97	109	121	133
(14)	2	14	26	38	50	62	74	86	98	110	122	134
(15)	3	15	27	39	51	63	75	87	99	111	123	135
(16)	4	16	28	40	52	64	76	88	100	112	124	136
(17)	5	17	29	41	53	65	77	89	101	113	125	137
(18)	6	18	30	42	54	66	78	90	102	114	126	138
(19)	7	19	31	43	55	67	79	91	103	115	127	139
(20)	8	20	32	44	56	68	80	92	104	116	128	140
(21)	9	21	33	45	57	69	81	93	105	117	129	141
(22)	10	22	34	46	58	70	82	94	106	118	130	142
(23)	11	23	35	47	59	71	83	95	107	119	131	143
(24)	12	24	36	48	60	72	84	96	108	120	132	144

	Rep. III											
(25)	1	14	27	40	57	70	83	96	101	114	127	140
(26)	2	13	28	39	58	69	84	95	102	113	128	139
(27)	3	16	25	38	59	72	81	94	103	116	125	138
(28)	4	15	26	37	60	71	82	93	104	115	126	137
(29)	5	18	31	44	49	62	75	88	105	118	131	144
(30)	6	17	32	43	50	61	76	87	106	117	132	143
(31)	7	20	29	42	51	64	73	86	107	120	129	142
(32)	8	19	30	41	52	63	74	85	108	119	130	141
(33)	9	22	35	48	53	66	79	92	97	110	123	136
(34)	10	21	36	47	54	65	80	91	98	109	124	135
(35)	11	24	33	46	55	68	77	90	99	112	121	134
(36)	12	23	34	45	56	67	78	89	100	111	122	133

Plan 10.9 (Continued)

12 × 12 quadruple lattice

	Rep. IV											
(37)	1	24	30	43	53	64	82	95	105	116	122	135
(38)	2	23	29	44	54	63	81	96	106	115	121	136
(39)	3	22	32	41	55	62	84	93	107	114	124	133
(40)	4	21	31	42	56	61	83	94	108	113	123	134
(41)	5	16	34	47	57	68	74	87	97	120	126	139
(42)	6	15	33	48	58	67	73	88	98	119	125	140
(43)	7	14	36	45	59	66	76	85	99	118	128	137
(44)	8	13	35	46	60	65	75	86	100	117	127	138
(45)	9	20	26	39	49	72	78	91	101	112	130	143
(46)	10	19	25	40	50	71	77	92	102	111	129	144
(47)	11	18	28	37	51	70	80	89	103	110	132	141
(48)	12	17	27	38	52	69	79	90	104	109	131	142

Plan 10.10

3 × 4 rectangular lattice

Block	Rep. X				Rep. Y				Rep. Z				
X1	1	2	3		Y1	4	7	10		Z1	6	8	12
X2	4	5	6		Y2	1	8	11		Z2	2	9	10
X3	7	8	9		Y3	2	5	12		Z3	3	4	11
X4	10	11	12		Y4	3	6	9		Z4	1	5	7

Plan 10.11

4 × 5 rectangular lattice

Block	Rep. X					Rep. Y					Rep. Z			
X1	1	2	3	4	Y1	5	9	13	17	Z1	8	11	15	18
X2	5	6	7	8	Y2	1	10	14	18	Z2	2	9	16	20
X3	9	10	11	12	Y3	2	6	15	19	Z3	4	7	14	17
X4	13	14	15	16	Y4	3	7	11	20	Z4	1	5	12	19
X5	17	18	19	20	Y5	4	8	12	16	Z5	3	6	10	13

Plan 10.12

5 × 6 rectangular lattice

Block	Rep. X						Rep. Y						Rep. Z						
X1	1	2	3	4	5		Y1	6	11	16	21	26		Z1	7	13	19	25	27
X2	6	7	8	9	10		Y2	1	12	17	22	27		Z2	5	14	16	23	29
X3	11	12	13	14	15		Y3	2	7	18	23	28		Z3	1	8	20	21	30
X4	16	17	18	19	20		Y4	3	8	13	24	29		Z4	2	9	15	22	26
X5	21	22	23	24	25		Y5	4	9	14	19	30		Z5	3	10	11	17	28
X6	26	27	28	29	30		Y6	5	10	15	20	25		Z6	4	6	12	18	24

Plan 10.13

 6×7 rectangular lattice

Block	Rep. X						Rep. Y						
X1	1	2	3	4	5	6	Y1	7	13	19	25	31	37
X2	7	8	9	10	11	12	Y2	1	14	20	26	32	38
X3	13	14	15	16	17	18	Y3	2	8	21	27	33	39
X4	19	20	21	22	23	24	Y4	3	9	15	28	34	40
X5	25	26	27	28	29	30	Y5	4	10	16	22	35	41
X6	31	32	33	34	35	36	Y6	5	11	17	23	29	42
X7	37	38	39	40	41	42	Y7	6	12	18	24	30	36

Rep. Z						
Z1	12	17	22	28	33	38
Z2	2	13	24	29	35	40
Z3	4	9	20	25	36	42
Z4	6	11	16	27	32	37
Z5	1	7	18	23	34	39
Z6	3	8	14	19	30	41
Z7	5	10	15	21	26	31

Plan 10.14

 7×8 rectangular lattice

Block	Rep. X							Rep. Y							
X1	1	2	3	4	5	6	7	Y1	8	15	22	29	36	43	50
X2	8	9	10	11	12	13	14	Y2	1	16	23	30	37	44	51
X3	15	16	17	18	19	20	21	Y3	2	9	24	31	38	45	52
X4	22	23	24	25	26	27	28	Y4	3	10	17	32	39	46	53
X5	29	30	31	32	33	34	35	Y5	4	11	18	25	40	47	54
X6	36	37	38	39	40	41	42	Y6	5	12	19	26	33	48	55
X7	43	44	45	46	47	48	49	Y7	6	13	20	27	34	41	56
X8	50	51	52	53	54	55	56	Y8	7	14	21	28	35	42	49

Rep. Z							
Z1	9	17	25	33	41	49	51
Z2	7	18	26	34	38	43	53
Z3	1	10	27	35	36	47	55
Z4	2	11	19	29	42	44	56
Z5	3	12	20	28	37	45	50
Z6	4	13	21	22	30	46	52
Z7	5	14	15	23	31	39	54
Z8	6	8	16	24	32	40	48

Plan 10.15

8 × 9 rectangular lattice

Block	Rep. X									Rep. Y							
X1	1	2	3	4	5	6	7	8	Y1	9	17	25	33	41	49	57	65
X2	9	10	11	12	13	14	15	16	Y2	1	18	26	34	42	50	58	66
X3	17	18	19	20	21	22	23	24	Y3	2	10	27	35	43	51	59	67
X4	25	26	27	28	29	30	31	32	Y4	3	11	19	36	44	52	60	68
X5	33	34	35	36	37	38	39	40	Y5	4	12	20	28	45	53	61	69
X6	41	42	43	44	45	46	47	48	Y6	5	13	21	29	37	54	62	70
X7	49	50	51	52	53	54	55	56	Y7	6	14	22	30	38	46	63	71
X8	57	58	59	60	61	62	63	64	Y8	7	15	23	31	39	47	55	72
X9	65	66	67	68	69	70	71	72	Y9	8	16	24	32	40	48	56	64

	Rep. Z							
Z1	16	23	30	37	45	52	59	66
Z2	2	17	32	39	46	54	61	68
Z3	4	11	26	33	48	55	63	70
Z4	6	13	20	35	42	49	64	72
Z5	8	15	22	29	44	51	58	65
Z6	1	9	24	31	38	53	60	67
Z7	3	10	18	25	40	47	62	69
Z8	5	12	19	27	34	41	56	71
Z9	7	14	21	28	36	43	50	57

Plan 10.16

9 × 10 rectangular lattice

Block	Rep. X								
X1	1	2	3	4	5	6	7	8	9
X2	10	11	12	13	14	15	16	17	18
X3	19	20	21	22	23	24	25	26	27
X4	28	29	30	31	32	33	34	35	36
X5	37	38	39	40	41	42	43	44	45
X6	46	47	48	49	50	51	52	53	54
X7	55	56	57	58	59	60	61	62	63
X8	64	65	66	67	68	69	70	71	72
X9	73	74	75	76	77	78	79	80	81
X10	82	83	84	85	86	87	88	89	90

	Rep. Y								
Y1	10	19	28	37	46	55	64	73	82
Y2	1	20	29	38	47	56	65	74	83
Y3	2	11	30	39	48	57	66	75	84
Y4	3	12	21	40	49	58	67	76	85
Y5	4	13	22	31	50	59	68	77	86
Y6	5	14	23	32	41	60	69	78	87
Y7	6	15	24	33	42	51	70	79	88
Y8	7	16	25	34	43	52	61	80	89
Y9	8	17	26	35	44	53	62	71	90
Y10	9	18	27	36	45	54	63	72	81

	Rep. Z								
Z1	11	21	31	41	51	61	71	81	83
Z2	9	22	32	42	52	62	64	76	84
Z3	1	12	33	43	53	63	68	73	87
Z4	2	13	23	44	54	55	65	79	89
Z5	3	14	24	34	46	56	72	75	90
Z6	4	15	25	35	45	57	67	74	82
Z7	5	16	26	36	37	47	66	77	85
Z8	6	17	27	28	38	48	58	78	86
Z9	7	18	19	29	39	49	59	69	88
Z10	8	10	20	30	40	50	60	70	80

CHAPTER 11

BALANCED INCOMPLETE BLOCKS

11.1 Description

The balanced lattices described in the last chapter are a particular group of a general class of designs known as balanced incomplete blocks. All these designs have the property that any pair of treatments appears together equally often within some block. Thus in plan 11.1 every pair of treatments appears together once in the same block, in plan 11.2 twice, and in plan 11.5 four times. This property insures that the same standard error may be used for comparing every pair of treatments; it also facilitates the statistical analysis, since any treatment total is adjusted in a single operation for all the blocks in which the treatment appears.

The construction of the designs presents interesting mathematical problems, (11.1), (11.2), (11.3). Although a design can be found for any number of treatments, t , and any size of block, k , most of these are of no interest for our purpose, since they require too many replications. The plans in this chapter have been restricted to those in which the number of replicates does not exceed 10.

The balanced lattices are the particular set of designs for which the relation $t = k^2$ holds. They also have the property that the blocks can be grouped in separate replications. The majority of the balanced incomplete block arrangements do not possess this property, which is possible only when t is a multiple of k . A few designs can be sorted into groups which comprize two or three replicates each.

In most plans, the block contains six or fewer units. Accordingly, the designs are adapted for experiments in which the appropriate size of block is small. They have been applied in greenhouse pot experiments, where the block is restricted to the width of the bench; in experiments on plant virus diseases, where the block consists of a small number of leaves on each plant; in experimental cookery where there is a limited number of stoves, in tests of mosquito repellents, reference (11.4), where the block consists of the two exposed arms of a subject; and in nutritional experiments where each child constitutes a block, 3 different foods being given during the 3 terms of the school year.

Examples from field experimental work occur in blueberry fertilizer trials where, owing to the uneven growth of different bushes, only a small number of them can be grouped into a homogeneous block. A similar consideration may limit the size of block in experiments on the control of fruit pests, where the tree constitutes the experimental unit.

A few types of factorial design—the 5×2 , 5×3 , 7×3 , and 7×4 —can be arranged in balanced incomplete blocks with the result that main effects and interactions are confounded to the same extent. It happens that in these cases there is no suitable method of confounding the interactions alone.

11.2 Comparisons with Other Designs

Much experimental material is of the type exemplified above, in that the natural grouping for use as a block accommodates only a small number of units. Where the number of treatments exceeds the number of units in a block, there are three alternatives to the use of a balanced incomplete block design. (1) A common treatment is placed in each block, so as to provide a basis for comparison between treatments which are in different blocks. (2) The treatments are divided into groups such that the number of treatments per group is equal to the number of units per block. The groups are compared in a split-plot arrangement (chapter 7). (3) Complete replicates are assembled by the grouping of units from more than one block, in order that a randomized block design may be used. The merits of the first two alternatives were discussed in section 9.5.

The relative efficiency of randomized blocks and balanced incomplete blocks has been discussed by Yates (11.5). Balanced incomplete block designs which are arranged in replications cannot be appreciably less accurate than randomized blocks. With balanced incomplete blocks which cannot be arranged in replications, a comparison with randomized blocks is not easy except by the use of uniformity data. Certain conclusions can be drawn from the nature of the experimental material. If there is some association among the members of different blocks, it may be possible to form replicates which are only slightly less homogeneous than the blocks. For example, suppose that in an animal experiment the litter is the natural block. If the responses of the animals are influenced by their genetic constitution and if replications are constructed from closely related litters, the experimental error may be only slightly greater within a replication than within a block. In this event randomized blocks may be the more accurate layout.

Sometimes the investigator has no information about any associations

between the groups which constitute the blocks; he knows merely that the experimental responses are likely to differ from group to group. In these circumstances Yates (11.5) has shown that incomplete blocks (with the recovery of inter-block information) are more accurate on the average except when the blocks are ineffective. There is the additional advantage that the block differences can be measured as a guide to the design of future experiments, whereas with replications composed of units from several blocks it would be difficult to estimate the block effects.

It is worth noting the *efficiency factor* (E) of any balanced incomplete block design whose use is under consideration. This factor, which is shown as a decimal fraction in table 11.3 (p. 327) and in the plans, is a lower limit to the efficiency of balanced incomplete blocks relative to randomized blocks. With 37 treatments in blocks of 9 units, for instance, the efficiency factor is 91%, so that the loss of efficiency could not exceed 9%. The loss equals 9% only in the rather unlikely situation in which the replications of 37 units are as uniform as the blocks of 9 units, yet the variation among blocks is large compared to that within blocks. With any reasonable prospect that the blocks are more homogeneous than the replicates, the incomplete block design is to be preferred in this case. On the other hand, with 7 treatments in blocks of 2 units, the efficiency factor is only 58%. In this case incomplete blocks will be more accurate only if there is a substantial reduction in error variation because of the reduction in block size.

11.3 Arrangement of Experimental Material

Units within the same incomplete block should be as homogeneous as possible. For those designs which can also be grouped in replications, it is advisable to place similar blocks, so far as these can be discerned, within the same replicate.

Certain of these designs can be rearranged to form an incomplete latin square (chapter 13) in which variation associated with two types of grouping can be eliminated from the experimental errors. The experimenter should consider whether advantage can be taken of this double control so as to reduce the error.

11.4 Randomization

The steps are:

1. Rearrange the blocks at random. (If the design is arranged in complete replications, the blocks are randomized only within each replication and the replications are kept separate.)

2. Randomize the positions of the treatment numbers within each block.
3. Assign treatments at random to the numbers in the plan.

11.5 Statistical Analysis

In order to present the shortest computational methods, it is advisable to divide the plans into five types, which are shown in the index at the end of this chapter.

11.51 Type I. Designs Arranged in Replications. We will analyze an example with 6 treatments in blocks of 2 (plan 11.3). The experiment was conducted by Dr. Pauline Paul at Iowa State College (11.6). Its object was to compare the effects of length of cold storage on the tenderness and flavor of beef roasts. Six periods of storage (0, 1, 2, 4, 9, and 18 days) were tested: these are denoted by treatment symbols 1, 2, 3, ..., 6, respectively.

Thirty roasts from the round of an animal were used. Four muscles each provided 6 roasts, while 3 muscles each provided 2 roasts. The roasts on any muscle group themselves naturally into pairs, since to each roast on the left side of an animal there corresponds another roast on the right side. From previous experience it was believed that the 2 roasts in any pair would give closely similar results. Variation among different pairs from the same muscle was expected to be somewhat larger, and variation among muscles to be still larger.

These opinions prompted the use of a design in blocks of 2, each block comprizing the left and right roasts in a pair. When grouping the blocks into replications, it was natural to put roasts from the same muscle into the same replicate. With the first 4 muscles a separate replicate could be made from each muscle. The remaining replication consisted of the 3 smaller muscles.

The plan and the scores for tenderness are given in table 11.1. Scoring was done by 4 judges, each marking on a scale from 0 to 10. The scores shown are their totals (out of 40), a high score indicating very tender beef.

The following symbols are used: $t = 6$ = the number of treatments; $k = 2$ = number of units per block; $r = 5$ = number of replications; $b = 15$ = number of blocks.

1. Find the block totals, B , the replicate totals, the grand total, G , and the treatment totals, T , placed below the plan.

TABLE 11.1 SCORES FOR TENDERNESS OF BEEF

Rep. I			Rep. II			Rep. III		
<i>B</i>			<i>B</i>			<i>B</i>		
(1) 7	(2) 17	24	(1) 17	(3) 27	44	(1) 10	(4) 25	35
(3) 26	(4) 25	51	(2) 23	(5) 27	50	(2) 26	(6) 37	63
(5) 33	(6) 29	62	(4) 29	(6) 30	59	(3) 24	(5) 26	50
<hr/>			<hr/>			<hr/>		
137			153			148		
<hr/>			<hr/>			<hr/>		
Rep. IV			Rep. V					
<i>B</i>			<i>B</i>					
(1) 25	(5) 40	65	(1) 11	(6) 27	38			
(2) 25	(4) 34	59	(2) 24	(3) 21	45			
(3) 34	(6) 32	66	(4) 26	(5) 32	58			
<hr/>			<hr/>					
190			141					
<hr/>			<hr/>					

Storage time (days)	Treatment no.	<i>T</i>	<i>B_t</i>	<i>Q</i>	<i>W</i>	<i>T</i> + μW	Adj. means
0	1	70	206	-66	19	71.8	14.4
1	2	115	241	-11	24	117.3	23.5
2	3	132	256	8	17	133.6	26.7
4	4	139	262	16	15	140.4	28.1
9	5	158	285	31	-24	155.7	31.1
18	6	155	288	22	-51	150.2	30.0
<hr/>		<i>G</i> = 769	1538	0	0	769.0	

2. For each treatment, calculate the following quantities:

B_t = total of all blocks in which the treatment appears

$$Q = kT - B_t = 2T - B_t$$

$$W = (t - k)T - (t - 1)B_t + (k - 1)G = 4T - 5B_t + G$$

The total of the Q 's and the W 's should each be zero.

3. The analysis of variance is as follows:

	d.f.	s.s.	m.s.
Replications	(<i>r</i> - 1)	4	298.5
Treatments (unadj.)	(<i>t</i> - 1)	5	1059.8
Blocks within reps. (adj.)	(<i>b</i> - <i>r</i>)	10	213.4
Intra-block error	(<i>tr</i> - <i>t</i> - <i>b</i> + 1)	10	77.3
Total	(<i>tr</i> - 1)	29	1649.0

The only sum of squares requiring special instructions is that for blocks within replications, adjusted for treatment effects. Since we do not know any simple way of finding this directly, it is obtained by a round-about method. Calculate in the usual way the *unadjusted* sum of squares for blocks within replications: this comes to 753.0. Then compute the sum of squares for treatments, adjusted for blocks, which is

$$\frac{(t-1) \sum Q^2}{ktr(k-1)} = \frac{5}{(2)(6)(5)(1)} \cdot [(66)^2 + (11)^2 + \cdots + (22)^2] = 520.2$$

The adjusted sum of squares for blocks is then given by the following combination of sums of squares

$$\begin{array}{rccccccc} \text{Blocks (unadj.)} & + & \text{treatments (adj.)} & - & \text{treatments (unadj.)} & & \\ 753.0 & + & 520.2 & - & 1059.8 & = & 213.4 \end{array}$$

4. Calculate the weighting factor

$$\begin{aligned} \mu &= \frac{r(E_b - E_c)}{rt(k-1)E_b + k(b-r-t+1)E_c} \\ &= \frac{5(21.34 - 7.73)}{(30)(21.34) + (10)(7.73)} = 0.09484 \end{aligned}$$

The adjusted treatment totals are

$$T + \mu W$$

and the means per unit are found on division by r .

5. The effective error variance per unit is estimated as

$$E_e[1 + (t-k)\mu] = 7.73[1 + (4)(0.0948)] = 10.66$$

We derive t -tests from this figure by the ordinary rules. For instance, the standard error of the difference between two adjusted totals is $\sqrt{2r(10.66)}$, or 10.32; for the difference between two adjusted *means* the standard error is $\sqrt{(2)(10.66)/r}$, or 2.06.

6. If an F -test is desired, calculate the sum of squares of the adjusted treatment totals. This figure, 4718.0, is divided by $r(t-1)$, or in this case 25, in order to find the mean square on a unit basis. The mean square, 188.7, is compared with the effective error m.s., 10.66, which is assigned $(tr - t - b + 1)$, or 10, d.f. The F -ratio, 188.7/10.66, is highly significant. The test is not exact, because it ignores the effects of errors in the weighting factor μ .

In an experiment of this type, interest centers on the shape of the response curve rather than on comparisons between pairs of treatments.

The adjusted means indicate the common "diminishing returns" curve. An exponential of the form $y = A - Be^{-ct}$, where t is the time of storage, fits the data satisfactorily. A parabolic regression on t , which is easier to compute, will be found to be inadequate. The conclusion is that storage up to about a week increases tenderness.

7. The gain in precision over randomized blocks is estimated as follows. By pooling the sums of squares for blocks within replications and intra-block error, we obtain an estimate of the error m.s. that would have applied with randomized blocks. This figure, 14.54, is compared with the effective error m.s., 10.66. The relative precision is estimated as $14.54/10.66$, or 136%. Since the randomized block design has 20 error d.f. as against 10 for the design actually used, this estimate should be reduced to 126% by the adjustment described in section 2.31.

11.52 Type II. Designs Arranged in Groups of Replications. Only two of the plans are of this type, plans 11.6 and 11.13, and in each case the group consists of *two* replications. (Plan 11.2 is also in this class, but is better analyzed as shown in section 11.55.) The analysis is very similar to that in the previous section. Two changes should be noted. If c is the number of groups, so that $c = r/2$ for the plans presented here, the degrees of freedom in the analysis of variance subdivide as shown below.

	d.f.	m.s.
Groups	$(c - 1)$	
Treatments (unadj.)	$(t - 1)$	
Blocks within groups (adj.)	$(b - c)$	E_b
Intra-block error	$(tr - t - b + 1)$	E_c
Total	$(tr - 1)$	

The second change is that the weighting factor μ becomes

$$\mu = \frac{(b - c)(E_b - E_c)}{t(k - 1)(b - c)E_b + (t - k)(b - c - t + 1)E_c}$$

11.53 Type III. Designs Not Arranged in Replications or Groups of Replications. Here again the method of section 11.51 may be followed except that no replication effects appear. The analysis of variance now reads as follows.

	d.f.	m.s.
Treatments (unadj.)	$(t - 1)$	
Blocks (adj.)	$(b - 1)$	E_b
Intra-block error	$(tr - t - b + 1)$	E_c
Total	$(tr - 1)$	

As before, the sum of squares for blocks (adjusted) is found by a subtraction process. The weighting factor becomes

$$\mu = \frac{(b-1)(E_b - E_e)}{t(k-1)(b-1)E_b + (t-k)(b-t)E_e}$$

Otherwise the method of section 11.51 applies. Of course, the gain in accuracy relative to randomized blocks cannot be estimated. A worked example with $t = 9$, $k = 4$, has been given by Yates (11.5, 11.7). His method of calculating the adjusted sum of squares for blocks is slightly different from ours, since he divides this sum of squares into two components so that the calculation becomes partially self-checking. Results should be identical by either method.

11.54 Type IV. Experiments with $t = b$. These designs, which cannot be arranged in replications, are covered by the computing instructions given in the previous section. However, the analysis simplifies slightly when $t = b$, because the sum of squares for blocks (adjusted) can be found directly. The data in table 11.2 are from an experiment on corn hybrids with $t = b = 13$, $r = k = 4$. For field experiments a design that can be laid out in separate replications is usually preferred. In this experiment, conducted in 1943 by the North Carolina Agricultural Experiment Station, it was desired to compare 10 hybrids, all potentially suitable for commercial use, at a number of places throughout the state, and at each place to test in addition 3 standard varieties that were adapted to that place. With 13 treatments no design that can be put into replications is available, and since 4 was a convenient number of replications, the design with $t = 13$, $r = k = 4$ was chosen. Randomized blocks might have been equally good. Incidentally, the plan used is different from that given in plan 11.22, a different method of construction having been employed. Both plans have the same structural properties.

1. Find the block totals, B , the treatment totals, T , and for each treatment find the total B_t of all blocks in which the treatment appears. Time is saved if T and B_t are found simultaneously. The B_t values should add to k times the grand total G .

2. For each treatment find

$$\begin{aligned} W &= (t-k)T - (t-1)B_t + (k-1)G \\ &= 9T - 12B_t + 3G \quad (\text{in this example}) \end{aligned}$$

The W 's should add to zero.

TABLE 11.2 PLAN AND YIELDS OF CORN (POUNDS PER PLOT)

Block									Block totals
1	(3)	25.3	(6)	19.9	(9)	29.0	(11)	24.6	98.8
2	(3)	23.0	(4)	19.8	(8)	33.3	(12)	22.7	98.8
3	(10)	16.2	(11)	19.3	(12)	31.7	(13)	26.6	93.8
4	(2)	27.3	(5)	27.0	(8)	35.6	(11)	17.4	107.3
5	(7)	23.4	(8)	30.5	(9)	30.8	(10)	32.4	117.1
6	(4)	30.6	(5)	32.4	(6)	27.2	(10)	32.8	123.0
7	(1)	34.7	(5)	31.1	(9)	25.7	(12)	30.5	122.0
8	(3)	34.4	(5)	32.4	(7)	33.3	(13)	36.9	137.0
9	(1)	38.2	(2)	32.9	(3)	37.3	(10)	31.3	139.7
10	(2)	28.7	(4)	30.7	(9)	26.9	(13)	35.3	121.6
11	(1)	36.6	(4)	31.1	(7)	31.1	(11)	28.4	127.2
12	(1)	31.8	(6)	33.7	(8)	27.8	(13)	41.1	134.4
13	(2)	30.3	(6)	31.5	(7)	39.3	(12)	26.7	127.8

1548.5

No.	T	B_t	W	$T + \mu W$
1	141.3	523.3	-362.4	136.7
2	119.2	496.4	-238.5	116.2
3	120.0	474.3	33.9	120.4
4	112.2	470.6	8.1	112.3
5	122.9	489.3	-120.0	121.4
6	112.3	484.0	-151.8	110.4
7	127.1	509.1	-319.8	123.0
8	127.2	457.6	299.1	131.0
9	112.4	459.5	143.1	114.2
10	112.7	473.6	-23.4	112.4
11	89.7	427.1	327.6	93.9
12	111.6	442.4	341.1	115.9
13	139.9	486.8	63.0	140.7
	1548.5	6194.0	0	1548.5

3. The analysis of variance is:

	d.f.		s.s.	m.s.
Treatments (unadj.)	$(t - 1)$	12	542.67	
Blocks (adj.)	$(b - 1)$	12	475.27	$39.61E_b$
Intra-block error	$(tr - 2t + 1)$	27	538.21	$19.93E_e$
Total	$(tr - 1)$	51	1556.15	

The sum of squares for blocks (adjusted) is found directly as the sum of squares of the W 's, divided by $tr(t - k)(k - 1)$, or in this case $(4)(13)(9)(3) = 1404$.

4. The weighting factor μ simplifies to

$$\mu = \frac{E_b - E_e}{t(k-1)E_b} = \frac{19.68}{(13)(3)(39.61)} = 0.0127$$

The adjusted treatment totals are the quantities $(T + \mu W)$.

5. As before, the effective error m.s. per unit is taken as

$$E_e[1 + (t - k)\mu] = 19.93[1 + (9)(0.0127)] = 22.2$$

Other features of the analysis are the same as in previous sections. Yates (11.5) presents an example with $t = 21$, $k = 5$.

11.55 Type V. Small Experiments. In certain of the smaller designs, the numbers of degrees of freedom in the mean squares for blocks and intra-block error are rather small. Consequently, the estimates of the relative weights assigned to inter- and intra-block comparisons are poor. In these cases, unless the whole plan has been repeated to secure extra replication, it is best to use the original method of analysis that Yates developed (11.8). This method adjusts the treatment means for differences between the blocks, but makes no use of inter-block information. The analysis is a little simpler than when inter-block information is used.

For purposes of illustration, this analysis will be applied to the example in the previous section, with $t = b = 13$, $r = k = 4$, though in practice inter-block information should be utilized in an experiment of this size. The data are presented in table 11.2.

1. Find the block totals, B , and the treatment totals, T . For each treatment find the total B_t over all blocks in which the treatment appears.

2. For each treatment compute

$$Q = kT - B_t = 4T - B_t$$

$$t' = m + \frac{t-1}{tr(k-1)} Q = m + \frac{12}{(13)(4)(3)} Q = m + 0.0769Q$$

where $m = 1548.5/52 = 29.78$ is the mean of the whole experiment. The t' values are the adjusted treatment means. The Q values should add to zero.

3. The analysis of variance is shown below.

	d.f.		s.s.	m.s.
Blocks (unadj.)	$(b-1)$	12	689.38	
Treatments (adj.)	$(t-1)$	12	328.55	27.38
Intra-block error	$(tr - t - b + 1)$	27	538.22	19.93 E_e
Total	$(tr-1)$	51	1556.15	

Note that the blocks sum of squares is *unadjusted*. It is found in the usual way as the sum of squares of deviations of the B 's, divided by k .

This rule applies even when the experiment is arranged in replications or groups of replications, though if the experimenter wishes, he may divide the blocks sum of squares into that due to replications (or groups) and that due to blocks within replications.

The adjusted sum of squares for treatments is

$$\frac{t-1}{rkt(k-1)} \sum Q^2 = \frac{12}{(4)(13)(4)(3)} [(41.9)^2 + (19.6)^2 + \cdots + (72.8)^2] \\ = 328.55$$

As usual, the intra-block error s.s. is found by subtraction.

4. For t -tests, the effective error variance is

$$E_e \left[1 + \frac{t-k}{t(k-1)} \right] = E_e \frac{k(t-1)}{t(k-1)} = \frac{(19.93)(12)(4)}{(13)(3)} = 24.5$$

For the variance of the difference between two adjusted treatment means, we multiply this quantity by $2/r$.

5. The analysis above provides an exact F -test of the adjusted treatment means. The F -ratio is $27.38/19.93$, with 12 and 27 d.f.

11.56 Missing Data. For reasons given in section 10.13, it is recommended that the analysis of incomplete data be carried out by inserting for the missing observations values which minimize the intra-block error s.s. In our notation the formula, first given by Cornish (11.9), is

$$x = \frac{tr(k-1)B + k(t-1)Q - (t-1)Q'}{(k-1)[tr(k-1) - k(t-1)]}$$

where B is the total of the block containing the missing observation, Q is the Q value ($= kT - B_t$) for the treatment that contains the missing value, and Q' is the sum of the Q values for all treatments that are in the block with the missing value.

Example 1. In the experiment on roast beef in section 11.51, suppose that treatment (5) in replication IV is absent. The first step is to find the block totals, the treatment totals, the B_t and the Q values as in the ordinary analysis. An x should be inserted for the missing observation. The data read as follows.

Treatment number	T	B_t	$Q = kT - B_t$
1	70	$166 + x$	$-26 - x$
2	115	241	-11
3	132	256	8
4	139	262	16
5	$118 + x$	$245 + x$	$-9 + x$
6	155	288	22
Totals	$729 + x$	$1458 + 2x$	0

Since the block with the missing value contains treatments (1) and (5)

$$Q' = -26 - 9 = -35$$

Hence, since B is 25,

$$\begin{aligned} x &= \frac{30B + 10Q - 5Q'}{20} = \frac{6B + 2Q - Q'}{4} \\ &= \frac{(6)(25) + (2)(-26) - (-35)}{4} = 42 \end{aligned}$$

On substituting this value in all places where an x occurs in the table above, we are ready to proceed with the analysis of variance.

Example 2. For an example with 2 missing observations, suppose that treatments (8) and (13) are missing from block 12 in the corn experiment in section 11.54. Denote the values to be inserted by x and y , respectively. To save space, the preliminary data are presented only for the treatments that occur in the same block as the missing observations; these are treatments (1), (6), (8), and (13).

Treatment number	T	B_t	$Q = 4T - B_t$
1	141.3	$454.4 + x + y$	$110.8 - x - y$
6	112.3	$415.1 + x + y$	$34.1 - x - y$
8	$99.4 + x$	$388.7 + x + y$	$8.9 + 3x - y$
13	$98.8 + y$	$417.9 + x + y$	$-22.7 - x + 3y$
			$131.1 = Q'$
$B = 65.5 + x + y$			

In the 3 replications in which it appears, the average for treatment (13) is 33: this will be chosen as the first approximation to y . Then for x we have

$$B = 65.5 + 33.0 = 98.5; \quad Q = 8.9 - 33 = -24.1; \quad Q' = 131.1$$

Since $t = 13$, $k = r = 4$,

$$\begin{aligned} x &= \frac{156B + 48Q - 12Q'}{324} = \frac{13B + 4Q - Q'}{27} \\ &= \frac{(13)(98.5) + (4)(-24.1) - (131.1)}{27} = 39.0 \end{aligned}$$

We now substitute $x = 39.0$ in the preliminary data and obtain a second approximation to y . For this we have

$$B = 65.5 + 39.0 = 104.5; \quad Q = -22.7 - 39.0 = -61.7$$

$$Q' = 131.1$$

Note that Q' , which does not involve x or y , remains fixed throughout the calculation. The second approximation to y is

$$y = \frac{(13)(104.5) + (4)(-61.7) - (131.1)}{27} = 36.3$$

The next 2 approximations will be found to be $x = 40.1$, $y = 36.7$, which may be used as the values to be inserted.

The rest of the analysis proceeds as usual; the intra-block error degrees of freedom are reduced by 1 for each missing observation. In the calculation of standard errors for t -tests of the adjusted treatment means, the method given in section 10.13 should be followed.

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TABLE 11.3 INDEX TO PLANS

t	k	r	b	λ †	E	Plan	Type
4	2	3	6	1	.67	11.1	V
	3	3	4	2	.89	*	V
5	2	4	10	1	.62	11.2	V
	3	6	10	3	.83	*	III
	4	4	5	3	.94	*	V
6	2	5	15	1	.60	11.3	I
	3	5	10	2	.80	11.4	III
	3	10	20	4	.80	11.5	I
	4	10	15	6	.90	11.6	II
	5	5	6	4	.96	*	V
7	2	6	21	1	.58	*	III
	3	3	7	1	.78	11.7	V
	4	4	7	2	.88	11.8	V
	6	6	7	5	.97	*	V

BALANCED INCOMPLETE BLOCKS

TABLE 11.3 INDEX TO PLANS (*Continued*)

t	k	r	b	λ^\dagger	E	Plan	Type
8	2	7	28	1	.57	11.9	I
	4	7	14	3	.86	11.10	I
	7	7	8	6	.98	*	V
9	2	8	36	1	.56	*	III
	4	8	18	3	.84	11.11	III
	5	10	18	5	.90	11.12	III
	6	8	12	5	.94	11.13	II
	8	8	9	7	.98	*	IV
10	2	9	45	1	.56	11.14	I
	3	9	30	2	.74	11.15	III
	4	6	15	2	.83	11.16	III
	5	9	18	4	.89	11.17	III
	6	9	15	5	.93	11.18	III
	9	9	10	8	.99	*	IV
11	2	10	55	1	.55	*	III
	5	5	11	2	.88	11.19	IV
	6	6	11	3	.92	11.20	IV
	10	10	11	9	.99	*	IV
13	3	6	26	1	.72	11.21	III
	4	4	13	1	.81	11.22	IV
	9	9	13	6	.96	11.23	IV
15	3	7	35	1	.71	11.24	I
	7	7	15	3	.92	11.25	IV
	8	8	15	4	.94	11.26	IV
16	6	6	16	2	.89	11.27	IV
	6	9	24	3	.89	11.28	III
	10	10	16	6	.96	11.29	IV
19	3	9	57	1	.70	11.30	III
	9	9	19	4	.94	11.31	IV
	10	10	19	5	.95	11.32	IV
21	3	10	70	1	.70	11.33	I
	5	5	21	1	.84	11.34	IV
	7	10	30	3	.90	11.35	III
25	4	8	50	1	.78	11.36	III
	9	9	25	3	.93	11.37	IV
28	4	9	63	1	.78	11.38	I
	7	9	36	2	.89	11.39	III
31	6	6	31	1	.86	11.40	IV
	10	10	31	3	.93	11.41	IV
37	9	9	37	2	.91	11.42	IV
41	5	10	82	1	.82	11.43	III
57	8	8	57	1	.89	11.44	IV
73	9	9	73	1	.90	11.45	IV
91	10	10	91	1	.91	11.46	IV

* These plans are constructed by forming all possible combinations of the t numbers in groups of size k . The number of blocks b serves as a check that no group has been missed.

† Number of times that two treatments appear together in the same block.

PLANS

Plan 11.1 $t = 4, k = 2, r = 3, b = 6, \lambda = 1, E = 0.67$, Type V

Block	Rep. I		Rep. II		Rep. III
(1)	<u>1 2</u>	(3)	<u>1 3</u>	(5)	<u>1 4</u>
(2)	<u>3 4</u>	(4)	<u>2 4</u>	(6)	<u>2 3</u>

Plan 11.2 $t = 5, k = 2, r = 4, b = 10, \lambda = 1, E = 0.62$, Type V

Block	Reps. I and II		Reps. III and IV
(1)	<u>1 2</u>	(6)	<u>1 4</u>
(2)	<u>3 4</u>	(7)	<u>2 3</u>
(3)	<u>2 5</u>	(8)	<u>3 5</u>
(4)	<u>1 3</u>	(9)	<u>1 5</u>
(5)	<u>4 5</u>	(10)	<u>2 4</u>

Plan 11.3 $t = 6, k = 2, r = 5, b = 15, \lambda = 1, E = 0.60$, Type I

Block	Rep. I		Rep. II		Rep. III		Rep. IV		Rep. V
(1)	<u>1 2</u>	(4)	<u>1 3</u>	(7)	<u>1 4</u>	(10)	<u>1 5</u>	(13)	<u>1 6</u>
(2)	<u>3 4</u>	(5)	<u>2 5</u>	(8)	<u>2 6</u>	(11)	<u>2 4</u>	(14)	<u>2 3</u>
(3)	<u>5 6</u>	(6)	<u>4 6</u>	(9)	<u>3 5</u>	(12)	<u>3 6</u>	(15)	<u>4 5</u>

Plan 11.4 $t = 6, k = 3, r = 5, b = 10, \lambda = 2, E = 0.80$, Type III

Block							
(1)	<u>1</u>	<u>2</u>	<u>5</u>	(6)	<u>2</u>	<u>3</u>	<u>4</u>
(2)	<u>1</u>	<u>2</u>	<u>6</u>	(7)	<u>2</u>	<u>3</u>	<u>5</u>
(3)	<u>1</u>	<u>3</u>	<u>4</u>	(8)	<u>2</u>	<u>4</u>	<u>6</u>
(4)	<u>1</u>	<u>3</u>	<u>6</u>	(9)	<u>3</u>	<u>5</u>	<u>6</u>
(5)	<u>1</u>	<u>4</u>	<u>5</u>	(10)	<u>4</u>	<u>5</u>	<u>6</u>

Plan 11.5 $t = 6, k = 3, r = 10, b = 20, \lambda = 4, E = 0.80$, Type I

Block	Rep. I		Rep. II		Rep. III		Rep. IV
(1)	<u>1 2 3</u>	(3)	<u>1 2 4</u>	(5)	<u>1 2 5</u>	(7)	<u>1 2 6</u>
(2)	<u>4 5 6</u>	(4)	<u>3 5 6</u>	(6)	<u>3 4 6</u>	(8)	<u>3 4 5</u>
	Rep. V		Rep. VI		Rep. VII		Rep. VIII
(9)	<u>1 3 4</u>	(11)	<u>1 3 5</u>	(13)	<u>1 3 6</u>	(15)	<u>1 4 5</u>
(10)	<u>2 5 6</u>	(12)	<u>2 4 6</u>	(14)	<u>2 4 5</u>	(16)	<u>2 3 6</u>
	Rep. IX		Rep. X				
(17)	<u>1 4 6</u>	(19)	<u>1 5 6</u>				
(18)	<u>2 3 5</u>	(20)	<u>2 3 4</u>				

Plan 11.6 $t = 6, k = 4, r = 10, b = 15, \lambda = 6, E = 0.90$, Type II

Block	Reps. I and II				Reps. III and IV				Reps. V and VI					
(1)	1	2	3	4	(4)	1	2	3	5	(7)	1	2	3	6
(2)	1	4	5	6	(5)	1	2	4	6	(8)	1	3	4	5
(3)	2	3	5	6	(6)	3	4	5	6	(9)	2	4	5	6
Reps. VII and VIII					Reps. IX and X									
(10)	1	2	4	5	(13)	1	2	5	6					
(11)	1	3	5	6	(14)	1	3	4	6					
(12)	2	3	4	6	(15)	2	3	4	5					

Plan 11.7 $t = 7, k = 3, r = 3, b = 7, \lambda = 1, E = 0.78$, Type V

Block															
(1)	<u>1</u>	<u>2</u>	<u>4</u>	(3)	<u>3</u>	<u>4</u>	<u>6</u>	(5)	<u>5</u>	<u>6</u>	<u>1</u>	(7)	<u>7</u>	<u>1</u>	<u>3</u>
(2)	<u>2</u>	<u>3</u>	<u>5</u>	(4)	<u>4</u>	<u>5</u>	<u>7</u>	(6)	<u>6</u>	<u>7</u>	<u>2</u>				

Plan 11.8 $t = 7, k = 4, r = 4, b = 7, \lambda = 2, E = 0.88$, Type V

Block														
(1)	3	5	6	7	(4)	1	2	3	6	(7)	2	4	5	6
(2)	1	4	6	7	(5)	2	3	4	7					
(3)	1	2	5	7	(6)	1	3	4	5					

Plan 11.9 $t = 8, k = 2, r = 7, b = 28, \lambda = 1, E = 0.57$, Type I

Block	Rep. I		Rep. II		Rep. III		Rep. IV	
(1)	1	2	(5)	1	3	(9)	1	4
(2)	3	4	(6)	2	8	(10)	2	7
(3)	5	6	(7)	4	5	(11)	3	6
(4)	7	8	(8)	6	7	(12)	5	8
	Rep. V		Rep. VI		Rep. VII			
(17)	1	6	(21)	1	7	(25)	1	8
(18)	2	4	(22)	2	6	(26)	2	5
(19)	3	8	(23)	3	5	(27)	3	7
(20)	5	7	(24)	4	8	(28)	4	6

Plan 11.10 $t = 8, k = 4, r = 7, b = 14, \lambda = 3, E = 0.86$, Type I

Block	Rep. I				Rep. II				Rep. III				Rep. IV						
(1)	1	2	3	4	(3)	1	2	7	8	(5)	1	3	6	8	(7)	1	4	6	7
(2)	5	6	7	8	(4)	3	4	5	6	(6)	2	4	5	7	(8)	2	3	5	8
Rep. V				Rep. VI				Rep. VII											
(9)	1	2	5	6	(11)	1	3	5	7	(13)	1	4	5	8					
(10)	3	4	7	8	(12)	2	4	6	8	(14)	2	3	6	7					

Plan 11.11 $t = 9, k = 4, r = 8, b = 18, \lambda = 3, E = 0.84$, Type III

Block

(1) <u>1 2 3 4</u>	(7) <u>1 4 8 9</u>	(13) <u>2 5 6 8</u>
(2) <u>1 2 5 6</u>	(8) <u>1 5 7 9</u>	(14) <u>3 5 8 9</u>
(3) <u>1 2 7 8</u>	(9) <u>2 3 8 9</u>	(15) <u>4 6 7 9</u>
(4) <u>1 3 5 7</u>	(10) <u>2 4 5 9</u>	(16) <u>3 4 5 6</u>
(5) <u>1 4 6 8</u>	(11) <u>2 6 7 9</u>	(17) <u>3 6 7 8</u>
(6) <u>1 3 6 9</u>	(12) <u>2 3 4 7</u>	(18) <u>4 5 7 8</u>

Plan 11.12 $t = 9, k = 5, r = 10, b = 18, \lambda = 5, E = 0.90$, Type III

Block

(1) <u>1 3 6 7 8</u>	(7) <u>1 3 4 7 9</u>	(13) <u>1 3 4 5 8</u>
(2) <u>2 3 4 6 8</u>	(8) <u>1 2 3 6 9</u>	(14) <u>1 2 4 6 7</u>
(3) <u>2 4 5 7 8</u>	(9) <u>1 2 3 5 8</u>	(15) <u>1 4 5 6 7</u>
(4) <u>5 6 7 8 9</u>	(10) <u>1 2 4 5 9</u>	(16) <u>2 3 5 7 9</u>
(5) <u>3 4 5 6 9</u>	(11) <u>3 4 7 8 9</u>	(17) <u>1 2 7 8 9</u>
(6) <u>2 4 6 8 9</u>	(12) <u>2 3 5 6 7</u>	(18) <u>1 5 6 8 9</u>

Plan 11.13 $t = 9, k = 6, r = 8, b = 12, \lambda = 5, E = 0.94$, Type II

Block	Reps. I and II	Reps. III and IV
(1)	<u>1 2 4 5 7 8</u>	(4) <u>1 2 5 6 7 9</u>
(2)	<u>2 3 5 6 8 9</u>	(5) <u>1 3 4 5 8 9</u>
(3)	<u>1 3 4 6 7 9</u>	(6) <u>2 3 4 6 7 8</u>
	Reps. V and VI	Reps. VII and VIII
(7)	<u>1 3 5 6 7 8</u>	(10) <u>4 5 6 7 8 9</u>
(8)	<u>1 2 4 6 8 9</u>	(11) <u>1 2 3 4 5 6</u>
(9)	<u>2 3 4 5 7 9</u>	(12) <u>1 2 3 7 8 9</u>

Plan 11.14 $t = 10, k = 2, r = 9, b = 45, \lambda = 1, E = 0.56$, Type I

Block	Rep. I	Rep. II	Rep. III	Rep. IV	Rep. V
(1)	<u>1 2</u>	(6) <u>1 3</u>	(11) <u>1 4</u>	(16) <u>1 5</u>	(21) <u>1 6</u>
(2)	<u>3 4</u>	(7) <u>2 7</u>	(12) <u>2 10</u>	(17) <u>2 8</u>	(22) <u>2 9</u>
(3)	<u>5 6</u>	(8) <u>4 8</u>	(13) <u>3 7</u>	(18) <u>3 10</u>	(23) <u>3 8</u>
(4)	<u>7 8</u>	(9) <u>5 9</u>	(14) <u>5 8</u>	(19) <u>4 9</u>	(24) <u>4 10</u>
(5)	<u>9 10</u>	(10) <u>6 10</u>	(15) <u>6 9</u>	(20) <u>6 7</u>	(25) <u>5 7</u>
	Rep. VI	Rep. VII	Rep. VIII	Rep. IX	
(26)	<u>1 7</u>	(31) <u>1 8</u>	(36) <u>1 9</u>	(41) <u>1 10</u>	
(27)	<u>2 6</u>	(32) <u>2 3</u>	(37) <u>2 4</u>	(42) <u>2 5</u>	
(28)	<u>3 9</u>	(33) <u>4 6</u>	(38) <u>3 5</u>	(43) <u>3 6</u>	
(29)	<u>4 5</u>	(34) <u>5 10</u>	(39) <u>6 8</u>	(44) <u>4 7</u>	
(30)	<u>8 10</u>	(35) <u>7 9</u>	(40) <u>7 10</u>	(45) <u>8 9</u>	

Plan 11.15 $t = 10, k = 3, r = 9, b = 30, \lambda = 2, E = 0.74$, Type III

Block

(1) <u>1 2 3</u>	(7) <u>1 7 9</u>	(13) <u>2 5 9</u>	(19) <u>3 5 6</u>	(25) <u>4 6 9</u>
(2) <u>1 2 4</u>	(8) <u>1 8 10</u>	(14) <u>2 6 7</u>	(20) <u>3 7 10</u>	(26) <u>4 7 8</u>
(3) <u>1 3 5</u>	(9) <u>1 9 10</u>	(15) <u>2 7 9</u>	(21) <u>3 8 9</u>	(27) <u>5 6 10</u>
(4) <u>1 4 6</u>	(10) <u>2 3 6</u>	(16) <u>2 8 10</u>	(22) <u>3 9 10</u>	(28) <u>5 7 8</u>
(5) <u>1 5 7</u>	(11) <u>2 4 10</u>	(17) <u>3 4 7</u>	(23) <u>4 5 9</u>	(29) <u>6 7 10</u>
(6) <u>1 6 8</u>	(12) <u>2 5 8</u>	(18) <u>3 4 8</u>	(24) <u>4 5 10</u>	(30) <u>6 8 9</u>

Plan 11.16 $t = 10, k = 4, r = 6, b = 15, \lambda = 2, E = 0.83$, Type III

Block

(1) <u>1 2 3 4</u>	(6) <u>1 6 8 10</u>	(11) <u>3 5 9 10</u>
(2) <u>1 2 5 6</u>	(7) <u>2 3 6 9</u>	(12) <u>3 6 7 10</u>
(3) <u>1 3 7 8</u>	(8) <u>2 4 7 10</u>	(13) <u>3 4 5 8</u>
(4) <u>1 4 9 10</u>	(9) <u>2 5 8 10</u>	(14) <u>4 5 6 7</u>
(5) <u>1 5 7 9</u>	(10) <u>2 7 8 9</u>	(15) <u>4 6 8 9</u>

Plan 11.17 $t = 10, k = 5, r = 9, b = 18, \lambda = 4, E = 0.89$, Type III

Block

(1) <u>1 2 3 4 5</u>	(7) <u>1 4 5 6 10</u>	(13) <u>2 5 6 8 10</u>
(2) <u>1 2 3 6 7</u>	(8) <u>1 4 8 9 10</u>	(14) <u>2 6 7 9 10</u>
(3) <u>1 2 4 6 9</u>	(9) <u>1 5 7 9 10</u>	(15) <u>3 4 6 7 10</u>
(4) <u>1 2 5 7 8</u>	(10) <u>2 3 4 8 10</u>	(16) <u>3 4 5 7 9</u>
(5) <u>1 3 6 8 9</u>	(11) <u>2 3 5 9 10</u>	(17) <u>3 5 6 8 9</u>
(6) <u>1 3 7 8 10</u>	(12) <u>2 4 7 8 9</u>	(18) <u>4 5 6 7 8</u>

Plan 11.18 $t = 10, k = 6, r = 9, b = 15, \lambda = 5, E = 0.93$, Type III

Block

(1) <u>1 2 4 5 8 9</u>	(6) <u>2 3 4 6 8 10</u>	(11) <u>1 4 5 7 8 10</u>
(2) <u>5 6 7 8 9 10</u>	(7) <u>1 2 6 7 9 10</u>	(12) <u>1 2 3 5 7 10</u>
(3) <u>2 4 5 6 9 10</u>	(8) <u>1 3 5 6 8 9</u>	(13) <u>2 3 5 6 7 8</u>
(4) <u>1 2 4 6 7 8</u>	(9) <u>1 2 3 8 9 10</u>	(14) <u>1 3 4 5 6 10</u>
(5) <u>3 4 7 8 9 10</u>	(10) <u>2 3 4 5 7 9</u>	(15) <u>1 3 4 6 7 9</u>

Plan 11.19 $t = 11, k = 5, r = 5, b = 11, \lambda = 2, E = 0.88$, Type IV

Block

(1) <u>1 2 3 5 8</u>	(7) <u>7 8 9 11 3</u>
(2) <u>2 3 4 6 9</u>	(8) <u>8 9 10 1 4</u>
(3) <u>3 4 5 7 10</u>	(9) <u>9 10 11 2 5</u>
(4) <u>4 5 6 8 11</u>	(10) <u>10 11 1 3 6</u>
(5) <u>5 6 7 9 1</u>	(11) <u>11 1 2 4 7</u>
(6) <u>6 7 8 10 2</u>	

Plan 11.20 $t = 11, k = 6, r = 6, b = 11, \lambda = 3, E = 0.92$, Type IV

Block

(1)	<u>4 6 7 9 10 11</u>	(7)	<u>1 2 4 5 6 10</u>
(2)	<u>1 5 7 8 10 11</u>	(8)	<u>2 3 5 6 7 11</u>
(3)	<u>1 2 6 8 9 11</u>	(9)	<u>1 3 4 6 7 8</u>
(4)	<u>1 2 3 7 9 10</u>	(10)	<u>2 4 5 7 8 9</u>
(5)	<u>2 3 4 8 10 11</u>	(11)	<u>3 5 6 8 9 10</u>
(6)	<u>1 3 4 5 9 11</u>		

Plan 11.21 $t = 13, k = 3, r = 6, b = 26, \lambda = 1, E = 0.72$, Type III

Block

(1)	<u>1 2 3</u>	(10)	<u>2 7 10</u>	(19)	<u>4 11 12</u>
(2)	<u>1 4 5</u>	(11)	<u>2 8 12</u>	(20)	<u>5 6 8</u>
(3)	<u>1 6 7</u>	(12)	<u>3 4 8</u>	(21)	<u>5 7 12</u>
(4)	<u>1 8 9</u>	(13)	<u>3 5 10</u>	(22)	<u>5 9 11</u>
(5)	<u>1 10 11</u>	(14)	<u>3 6 12</u>	(23)	<u>6 9 13</u>
(6)	<u>1 12 13</u>	(15)	<u>3 7 9</u>	(24)	<u>7 8 11</u>
(7)	<u>2 4 9</u>	(16)	<u>3 11 13</u>	(25)	<u>8 10 13</u>
(8)	<u>2 5 13</u>	(17)	<u>4 6 10</u>	(26)	<u>9 10 12</u>
(9)	<u>2 6 11</u>	(18)	<u>4 7 13</u>		

Plan 11.22 $t = 13, k = 4, r = 4, b = 13, \lambda = 1, E = 0.81$, Type IV

Block

(1)	<u>1 2 4 10</u>	(6)	<u>6 7 9 2</u>	(11)	<u>11 12 1 7</u>
(2)	<u>2 3 5 11</u>	(7)	<u>7 8 10 3</u>	(12)	<u>12 13 2 8</u>
(3)	<u>3 4 6 12</u>	(8)	<u>8 9 11 4</u>	(13)	<u>13 1 3 9</u>
(4)	<u>4 5 7 13</u>	(9)	<u>9 10 12 5</u>		
(5)	<u>5 6 8 1</u>	(10)	<u>10 11 13 6</u>		

Plan 11.23 $t = 13, k = 9, r = 9, b = 13, \lambda = 6, E = 0.96$, Type IV

Block

(1)	<u>3 5 6 7 8 9 11 12 13</u>	(8)	<u>1 2 3 5 6 7 10 12 13</u>
(2)	<u>1 4 6 7 8 9 10 12 13</u>	(9)	<u>1 2 3 4 6 7 8 11 13</u>
(3)	<u>1 2 5 7 8 9 10 11 13</u>	(10)	<u>1 2 3 4 5 7 8 9 12</u>
(4)	<u>1 2 3 6 8 9 10 11 12</u>	(11)	<u>2 3 4 5 6 8 9 10 13</u>
(5)	<u>2 3 4 7 9 10 11 12 13</u>	(12)	<u>1 3 4 5 6 7 9 10 11</u>
(6)	<u>1 3 4 5 8 10 11 12 13</u>	(13)	<u>2 4 5 6 7 8 10 11 12</u>
(7)	<u>1 2 4 5 6 9 11 12 13</u>		

Plan 11.24 $t = 15, k = 3, r = 7, b = 35, \lambda = 1, E = 0.71$, Type I

Block	Rep. I		Rep. II		Rep. III		Rep. IV
(1)	<u>1 2 3</u>	(6)	<u>1 4 5</u>	(11)	<u>1 6 7</u>	(16)	<u>1 8 9</u>
(2)	<u>4 8 12</u>	(7)	<u>2 8 10</u>	(12)	<u>2 9 11</u>	(17)	<u>2 13 15</u>
(3)	<u>5 10 15</u>	(8)	<u>3 13 14</u>	(13)	<u>3 12 15</u>	(18)	<u>3 4 7</u>
(4)	<u>6 11 13</u>	(9)	<u>6 9 15</u>	(14)	<u>4 10 14</u>	(19)	<u>5 11 14</u>
(5)	<u>7 9 14</u>	(10)	<u>7 11 12</u>	(15)	<u>5 8 13</u>	(20)	<u>6 10 12</u>
	Rep. V		Rep. VI		Rep. VII		
(21)	<u>1 10 11</u>	(26)	<u>1 12 13</u>	(31)	<u>1 14 15</u>		
(22)	<u>2 12 14</u>	(27)	<u>2 5 7</u>	(32)	<u>2 4 6</u>		
(23)	<u>3 5 6</u>	(28)	<u>3 9 10</u>	(33)	<u>3 8 11</u>		
(24)	<u>4 9 13</u>	(29)	<u>4 11 15</u>	(34)	<u>5 9 12</u>		
(25)	<u>7 8 15</u>	(30)	<u>6 8 14</u>	(35)	<u>7 10 13</u>		

Plan 11.25 $t = 15, k = 7, r = 7, b = 15, \lambda = 3, E = 0.92$, Type IV

Block

(1)	<u>1 2 3 4 5 6 7</u>	(9)	<u>2 4 6 9 11 12 14</u>
(2)	<u>1 2 3 8 9 10 11</u>	(10)	<u>2 5 7 8 10 12 14</u>
(3)	<u>1 2 3 12 13 14 15</u>	(11)	<u>2 5 7 9 11 13 15</u>
(4)	<u>1 4 5 8 9 14 15</u>	(12)	<u>3 4 7 8 11 13 14</u>
(5)	<u>1 4 5 10 11 12 13</u>	(13)	<u>3 4 7 9 10 12 15</u>
(6)	<u>1 6 7 8 9 12 13</u>	(14)	<u>3 5 6 8 11 12 15</u>
(7)	<u>1 6 7 10 11 14 15</u>	(15)	<u>3 5 6 9 10 13 14</u>
(8)	<u>2 4 6 8 10 13 15</u>		

Plan 11.26 $t = 15, k = 8, r = 8, b = 15, \lambda = 4, E = 0.94$, Type IV

Block

(1)	<u>8 9 10 11 12 13 14 15</u>	(9)	<u>1 3 5 7 8 10 13 15</u>
(2)	<u>4 5 6 7 12 13 14 15</u>	(10)	<u>1 3 4 6 9 11 13 15</u>
(3)	<u>4 5 6 7 8 9 10 11</u>	(11)	<u>1 3 4 6 8 10 12 14</u>
(4)	<u>2 3 6 7 10 11 12 13</u>	(12)	<u>1 2 5 6 9 10 12 15</u>
(5)	<u>2 3 6 7 8 9 14 15</u>	(13)	<u>1 2 5 6 8 11 13 14</u>
(6)	<u>2 3 4 5 10 11 14 15</u>	(14)	<u>1 2 4 7 9 10 13 14</u>
(7)	<u>2 3 4 5 8 9 12 13</u>	(15)	<u>1 2 4 7 8 11 12 15</u>
(8)	<u>1 3 5 7 9 11 12 14</u>		

Plan 11.27 $t = 16, k = 6, r = 6, b = 16, \lambda = 2, E = 0.89$, Type IV

Block

(1)	13	14	15	4	8	12
(2)	6	7	8	1	9	13
(3)	9	10	11	4	8	16
(4)	5	6	8	3	11	15
(5)	9	11	12	2	6	14
(6)	13	15	16	2	6	10
(7)	9	10	12	3	7	15
(8)	10	11	12	1	5	13

(9)	5	7	8	2	10	14
(10)	1	2	3	8	12	16
(11)	1	3	4	6	10	14
(12)	13	14	16	3	7	11
(13)	1	2	4	7	11	15
(14)	5	6	7	4	12	16
(15)	2	3	4	5	9	13
(16)	14	15	16	1	5	9

Plan 11.28 $t = 16, k = 6, r = 9, b = 24, \lambda = 3, E = 0.89$, Type III

Block

(1)	1	2	5	6	11	12
(2)	1	2	7	8	13	14
(3)	1	2	9	10	15	16
(4)	1	3	5	7	10	12
(5)	1	3	6	8	13	15
(6)	1	3	9	11	14	16
(7)	1	4	5	8	10	11
(8)	1	4	6	7	13	16
(9)	1	4	9	12	14	15
(10)	2	3	5	8	14	15
(11)	2	3	6	7	9	12
(12)	2	3	10	11	13	16

(13)	2	4	5	7	14	16
(14)	2	4	6	8	9	11
(15)	2	4	10	12	13	15
(16)	3	4	5	6	15	16
(17)	3	4	7	8	9	10
(18)	3	4	11	12	13	14
(19)	5	6	9	10	13	14
(20)	5	7	9	11	13	15
(21)	5	8	9	12	13	16
(22)	6	7	10	11	14	15
(23)	6	8	10	12	14	16
(24)	7	8	11	12	15	16

Plan 11.29 $t = 16, k = 10, r = 10, b = 16, \lambda = 6, E = 0.96$, Type IV

Block

(1)	1	2	3	5	6	7	9	10	11	16
(2)	2	3	4	5	10	11	12	14	15	16
(3)	1	2	3	5	6	7	12	13	14	15
(4)	1	2	4	7	9	10	12	13	14	16
(5)	1	3	4	5	7	8	10	13	15	16
(6)	1	3	4	5	7	8	9	11	12	14
(7)	1	2	4	5	6	8	11	13	14	16
(8)	2	3	4	6	7	8	9	14	15	16

(9)	1	3	4	6	9	11	12	13	15	16
(10)	4	5	6	7	9	10	11	13	14	15
(11)	2	5	7	8	9	11	12	13	15	16
(12)	1	2	4	5	6	8	9	10	12	15
(13)	3	5	6	8	9	10	12	13	14	16
(14)	1	2	3	8	9	10	11	13	14	15
(15)	1	6	7	8	10	11	12	14	15	16
(16)	2	3	4	6	7	8	10	11	12	13

Plan 11.30 $t = 19, k = 3, r = 9, b = 57, \lambda = 1, E = 0.70$, Type III

Block

(1) <u>1 2 19</u>	(16) <u>2 9 11</u>	(30) <u>4 13 14</u>	(44) <u>8 14 19</u>
(2) <u>1 3 6</u>	(17) <u>2 10 18</u>	(31) <u>5 6 11</u>	(45) <u>8 15 18</u>
(3) <u>1 4 18</u>	(18) <u>3 4 15</u>	(32) <u>5 7 10</u>	(46) <u>8 16 17</u>
(4) <u>1 5 16</u>	(19) <u>3 5 14</u>	(33) <u>5 8 9</u>	(47) <u>9 13 19</u>
(5) <u>1 7 14</u>	(20) <u>3 7 12</u>	(34) <u>5 12 17</u>	(48) <u>9 14 18</u>
(6) <u>1 8 13</u>	(21) <u>3 8 11</u>	(35) <u>5 13 18</u>	(49) <u>10 12 19</u>
(7) <u>1 9 12</u>	(22) <u>3 10 13</u>	(36) <u>6 7 8</u>	(50) <u>10 14 17</u>
(8) <u>1 10 11</u>	(23) <u>3 9 16</u>	(37) <u>6 9 10</u>	(51) <u>10 15 16</u>
(9) <u>1 15 17</u>	(24) <u>3 18 19</u>	(38) <u>6 13 15</u>	(52) <u>11 12 18</u>
(10) <u>2 3 17</u>	(25) <u>4 5 19</u>	(39) <u>6 16 19</u>	(53) <u>11 13 17</u>
(11) <u>2 4 16</u>	(26) <u>4 6 12</u>	(40) <u>6 17 18</u>	(54) <u>11 14 16</u>
(12) <u>2 5 15</u>	(27) <u>4 7 11</u>	(41) <u>7 9 15</u>	(55) <u>11 15 19</u>
(13) <u>2 6 14</u>	(28) <u>4 8 10</u>	(42) <u>7 16 18</u>	(56) <u>12 13 16</u>
(14) <u>2 7 13</u>	(29) <u>4 9 17</u>	(43) <u>7 17 19</u>	(57) <u>12 14 15</u>
(15) <u>2 8 12</u>			

Plan 11.31 $t = 19, k = 9, r = 9, b = 19, \lambda = 4, E = 0.94$, Type IV

Block

(1) <u>1 2 3 4 5 6 7 8 9</u>	(11) <u>2 3 5 6 11 13 16 17 18</u>
(2) <u>1 2 4 5 10 13 14 17 19</u>	(12) <u>2 3 8 9 10 12 17 18 19</u>
(3) <u>1 2 7 8 11 12 14 16 17</u>	(13) <u>2 4 6 8 10 14 15 16 18</u>
(4) <u>1 2 6 9 12 13 15 16 19</u>	(14) <u>2 5 7 9 11 14 15 18 19</u>
(5) <u>1 3 4 6 11 12 14 18 19</u>	(15) <u>3 4 5 9 12 14 15 16 17</u>
(6) <u>1 3 7 9 10 13 14 16 18</u>	(16) <u>3 6 7 8 13 14 15 17 19</u>
(7) <u>1 3 5 8 10 11 15 16 19</u>	(17) <u>4 5 7 8 12 13 16 18 19</u>
(8) <u>1 4 8 9 11 13 15 17 18</u>	(18) <u>4 6 7 9 10 11 16 17 19</u>
(9) <u>1 5 6 7 10 12 15 17 18</u>	(19) <u>5 6 8 9 10 11 12 13 14</u>
(10) <u>2 3 4 7 10 11 12 13 15</u>	

Plan 11.32 $t = 19, k = 10, r = 10, b = 19, \lambda = 5, E = 0.95$, Type IV

Block

(1)	<u>10 11 12 13 14 15 16 17 18 19</u>	(11)	<u>1 4 7 8 9 10 12 14 15 19</u>
(2)	<u>3 6 7 8 9 11 12 15 16 18</u>	(12)	<u>1 4 5 6 7 11 13 14 15 16</u>
(3)	<u>3 4 5 6 9 10 13 15 18 19</u>	(13)	<u>1 3 5 7 9 11 12 13 17 19</u>
(4)	<u>3 4 5 7 8 10 11 14 17 18</u>	(14)	<u>1 3 4 6 8 10 12 13 16 17</u>
(5)	<u>2 5 7 8 9 10 13 15 16 17</u>	(15)	<u>1 2 6 7 8 10 11 13 18 19</u>
(6)	<u>2 4 5 6 8 11 12 15 17 19</u>	(16)	<u>1 2 4 5 9 10 11 12 16 18</u>
(7)	<u>2 4 6 7 9 12 13 14 17 18</u>	(17)	<u>1 2 3 6 9 10 11 14 15 17</u>
(8)	<u>2 3 5 6 7 10 12 14 16 19</u>	(18)	<u>1 2 3 5 8 12 13 14 15 18</u>
(9)	<u>2 3 4 8 9 11 13 14 16 19</u>	(19)	<u>1 2 3 4 7 15 16 17 18 19</u>
(10)	<u>1 5 6 8 9 14 16 17 18 19</u>		

Plan 11.33 $t = 21, k = 3, r = 10, b = 70, \lambda = 1, E = 0.70$, Type I

Block Rep. I	Rep. II	Rep. III	Rep. IV.	Rep. V
(1) <u>1 2 3</u>	(8) <u>1 4 15</u>	(15) <u>1 5 17</u>	(22) <u>1 6 9</u>	(29) <u>1 7 21</u>
(2) <u>4 5 6</u>	(9) <u>2 5 11</u>	(16) <u>2 4 14</u>	(23) <u>2 7 16</u>	(30) <u>2 13 17</u>
(3) <u>7 8 9</u>	(10) <u>3 9 16</u>	(17) <u>3 7 11</u>	(24) <u>3 8 21</u>	(31) <u>3 10 18</u>
(4) <u>10 11 12</u>	(11) <u>6 17 20</u>	(18) <u>6 10 19</u>	(25) <u>4 17 19</u>	(32) <u>4 8 11</u>
(5) <u>13 14 15</u>	(12) <u>7 12 19</u>	(19) <u>8 16 20</u>	(26) <u>5 10 13</u>	(33) <u>5 16 19</u>
(6) <u>16 17 18</u>	(13) <u>8 13 18</u>	(20) <u>9 15 18</u>	(27) <u>11 15 20</u>	(34) <u>6 12 15</u>
(7) <u>19 20 21</u>	(14) <u>10 14 21</u>	(21) <u>12 13 21</u>	(28) <u>12 14 18</u>	(35) <u>9 14 20</u>
Rep. VI	Rep. VII	Rep. VIII	Rep. IX	Rep. X
(36) <u>1 8 10</u>	(43) <u>1 11 18</u>	(50) <u>1 12 20</u>	(57) <u>1 13 19</u>	(64) <u>1 14 16</u>
(37) <u>2 18 19</u>	(44) <u>2 10 20</u>	(51) <u>2 6 8</u>	(58) <u>2 9 12</u>	(65) <u>2 15 21</u>
(38) <u>3 15 17</u>	(45) <u>3 5 12</u>	(52) <u>3 14 19</u>	(59) <u>3 4 20</u>	(66) <u>3 6 13</u>
(39) <u>4 12 16</u>	(46) <u>4 9 13</u>	(53) <u>4 18 21</u>	(60) <u>5 8 14</u>	(67) <u>4 7 10</u>
(40) <u>5 9 21</u>	(47) <u>6 16 21</u>	(54) <u>5 7 15</u>	(61) <u>6 7 18</u>	(68) <u>5 18 20</u>
(41) <u>6 11 14</u>	(48) <u>7 14 17</u>	(55) <u>9 10 17</u>	(62) <u>10 15 16</u>	(69) <u>8 12 17</u>
(42) <u>7 13 20</u>	(49) <u>8 15 19</u>	(56) <u>11 13 16</u>	(63) <u>11 17 21</u>	(70) <u>9 11 19</u>

Plan 11.34 $t = 21, k = 5, r = 5, b = 21, \lambda = 1, E = 0.84$, Type IV

Block

(1) <u>1 2 3 4 17</u>	(8) <u>4 8 12 16 18</u>	(15) <u>3 5 10 16 20</u>
(2) <u>5 6 7 8 17</u>	(9) <u>1 6 11 16 19</u>	(16) <u>4 6 9 15 20</u>
(3) <u>9 10 11 12 17</u>	(10) <u>2 5 12 15 19</u>	(17) <u>1 8 10 15 21</u>
(4) <u>13 14 15 16 17</u>	(11) <u>3 8 9 14 19</u>	(18) <u>2 7 9 16 21</u>
(5) <u>1 5 9 13 18</u>	(12) <u>4 7 10 13 19</u>	(19) <u>3 6 12 13 21</u>
(6) <u>2 6 10 14 18</u>	(13) <u>1 7 12 14 20</u>	(20) <u>4 5 11 14 21</u>
(7) <u>3 7 11 15 18</u>	(14) <u>2 8 11 13 20</u>	(21) <u>17 18 19 20 21</u>

Plan 11.35 $t = 21, k = 7, r = 10, b = 30, \lambda = 3, E = 0.90$, Type III

Block

(1)	2 5 10 11 17 19 20	(16)	2 7 10 13 18 20 21
(2)	3 6 11 12 18 20 21	(17)	3 1 11 14 19 21 15
(3)	4 7 12 13 19 21 15	(18)	4 2 12 8 20 15 16
(4)	5 1 13 14 20 15 16	(19)	5 3 13 9 21 16 17
(5)	6 2 14 8 21 16 17	(20)	6 4 14 10 15 17 18
(6)	7 3 8 9 15 17 18	(21)	7 5 8 11 16 18 19
(7)	1 4 9 10 16 18 19	(22)	1 2 4 8 9 11 21
(8)	3 4 8 13 17 19 20	(23)	2 3 5 9 10 12 15
(9)	4 5 9 14 18 20 21	(24)	3 4 6 10 11 13 16
(10)	5 6 10 8 19 21 15	(25)	4 5 7 11 12 14 17
(11)	6 7 11 9 20 15 16	(26)	5 6 1 12 13 8 18
(12)	7 1 12 10 21 16 17	(27)	6 7 2 13 14 9 19
(13)	1 2 13 11 15 17 18	(28)	7 1 3 14 8 10 20
(14)	2 3 14 12 16 18 19	(29)	1 2 3 4 5 6 7
(15)	1 6 9 12 17 19 20	(30)	8 9 10 11 12 13 14

Plan 11.36 $t = 25, k = 4, r = 8, b = 50, \lambda = 1, E = 0.78$, Type III

Block

(1)	25 24 8 16	(18)	23 2 21 15	(35)	3 5 2 16
(2)	1 3 24 14	(19)	2 9 14 13	(36)	17 15 9 20
(3)	19 10 3 9	(20)	9 22 11 8	(37)	16 13 22 1
(4)	7 23 10 5	(21)	22 7 19 4	(38)	20 8 7 3
(5)	13 8 23 6	(22)	8 4 2 12	(39)	10 6 4 14
(6)	24 18 4 20	(23)	4 5 9 25	(40)	21 12 5 11
(7)	18 6 5 1	(24)	5 15 22 24	(41)	14 25 15 19
(8)	6 12 15 3	(25)	15 13 7 18	(42)	11 24 13 10
(9)	12 25 13 17	(26)	25 2 10 18	(43)	23 16 18 9
(10)	3 17 18 11	(27)	1 4 23 17	(44)	2 20 6 22
(11)	17 16 6 19	(28)	19 18 8 21	(45)	9 1 12 7
(12)	16 20 12 10	(29)	7 17 24 2	(46)	22 3 25 23
(13)	20 1 25 21	(30)	13 21 3 4	(47)	8 14 17 5
(14)	10 21 17 22	(31)	24 9 21 6	(48)	4 11 16 15
(15)	21 14 16 7	(32)	18 22 14 12	(49)	5 19 20 13
(16)	14 11 20 23	(33)	6 7 11 25	(50)	15 10 1 8
(17)	11 19 1 2	(34)	12 23 19 24		

Plan 11.37 $t = 25, k = 9, r = 9, b = 25, \lambda = 3, E = 0.93$, Type IV

Block

(1)	1 2 3 4 5 6 7 8 9	(14)	2 6 7 10 12 18 20 23 25
(2)	1 2 3 14 15 18 19 22 23	(15)	2 6 7 11 13 19 21 22 24
(3)	1 2 3 16 17 20 21 24 25	(16)	3 4 8 10 13 18 21 23 24
(4)	1 4 7 10 11 14 15 20 21	(17)	3 4 8 11 12 19 20 22 25
(5)	1 4 7 12 13 16 17 18 19	(18)	3 5 7 10 13 15 16 22 25
(6)	1 5 9 10 11 18 19 24 25	(19)	3 5 7 11 12 14 17 23 24
(7)	1 5 9 12 13 20 21 22 23	(20)	3 6 9 10 13 14 17 19 20
(8)	1 6 8 10 11 16 17 22 23	(21)	3 6 9 11 12 15 16 18 21
(9)	1 6 8 12 13 14 15 24 25	(22)	4 5 6 14 16 18 20 22 24
(10)	2 4 9 10 12 15 17 22 24	(23)	4 5 6 15 17 19 21 23 25
(11)	2 4 9 11 13 14 16 23 25	(24)	7 8 9 14 17 18 21 22 25
(12)	2 5 8 10 12 14 16 19 21	(25)	7 8 9 15 16 19 20 23 24
(13)	2 5 8 11 13 15 17 18 20		

Plan 11.38 $t = 28, k = 4, r = 9, b = 63, \lambda = 1, E = 0.78$, Type I

Block

Block	Rep. I				Rep. II				Rep. III					
(1)	28	1	10	19	(8)	28	2	11	20	(15)	28	3	12	21
(2)	2	9	13	16	(9)	3	1	14	17	(16)	4	2	15	18
(3)	3	8	11	18	(10)	4	9	12	10	(17)	5	1	13	11
(4)	4	7	23	24	(11)	5	8	24	25	(18)	6	9	25	26
(5)	5	6	20	27	(12)	6	7	21	19	(19)	7	8	22	20
(6)	12	17	22	25	(13)	13	18	23	26	(20)	14	10	24	27
(7)	14	15	21	26	(14)	15	16	22	27	(21)	16	17	23	19
Rep. IV				Rep. V				Rep. VI						
(22)	28	4	13	22	(29)	28	5	14	23	(36)	28	6	15	24
(23)	5	3	16	10	(30)	6	4	17	11	(37)	7	5	18	12
(24)	6	2	14	12	(31)	7	3	15	13	(38)	8	4	16	14
(25)	7	1	26	27	(32)	8	2	27	19	(39)	9	3	19	20
(26)	8	9	23	21	(33)	9	1	24	22	(40)	1	2	25	23
(27)	15	11	25	19	(34)	16	12	26	20	(41)	17	13	27	21
(28)	17	18	24	20	(35)	18	10	25	21	(42)	10	11	26	22
Rep. VII				Rep. VIII				Rep. IX						
(43)	28	7	16	25	(50)	28	8	17	26	(57)	28	9	18	27
(44)	8	6	10	13	(51)	9	7	11	14	(58)	1	8	12	15
(45)	9	5	17	15	(52)	1	6	18	16	(59)	2	7	10	17
(46)	1	4	20	21	(53)	2	5	21	22	(60)	3	6	22	23
(47)	2	3	26	24	(54)	3	4	27	25	(61)	4	5	19	26
(48)	18	14	19	22	(55)	10	15	20	23	(62)	11	16	21	24
(49)	11	12	27	23	(56)	12	13	19	24	(63)	13	14	20	25

Plan 11.39 $t = 28, k = 7, r = 9, b = 36, \lambda = 2, E = 0.89$, Type III

Block

(1)	4 7 8 9 14 23 28	(19)	4 8 11 17 19 21 25
(2)	1 5 9 10 11 15 24	(20)	1 13 14 18 23 25 26
(3)	6 8 13 15 16 18 21	(21)	2 4 5 6 16 22 23
(4)	7 12 13 17 22 24 25	(22)	3 4 10 11 12 14 18
(5)	4 10 16 17 20 26 27	(23)	1 9 14 16 17 19 22
(6)	2 11 18 19 22 26 28	(24)	1 2 4 13 20 24 28
(7)	1 3 6 12 19 23 27	(25)	3 5 8 17 23 24 26
(8)	2 3 5 14 20 21 25	(26)	5 6 7 10 19 25 28
(9)	1 2 8 10 12 16 25	(27)	1 6 7 8 11 20 26
(10)	2 3 6 9 11 13 17	(28)	9 10 13 19 20 21 23
(11)	4 5 12 13 15 19 26	(29)	2 8 14 15 19 24 27
(12)	3 7 16 18 19 20 24	(30)	3 9 15 16 25 26 28
(13)	6 10 14 21 22 24 26	(31)	5 8 9 12 18 20 22
(14)	11 15 20 22 23 25 27	(32)	11 12 16 21 23 24 28
(15)	1 5 17 18 21 27 28	(33)	1 3 4 7 15 21 22
(16)	2 7 9 12 21 26 27	(34)	5 7 11 13 14 16 27
(17)	3 8 10 13 22 27 28	(35)	4 6 9 18 24 25 27
(18)	6 12 14 15 17 20 28	(36)	2 7 10 15 17 18 23

Plan 11.40 $t = 31, k = 6, r = 6, b = 31, \lambda = 1, E = 0.86$, Type IV

Block

(1)	1 6 11 16 21 26	(12)	6 2 23 19 15 28	(22)	11 2 18 9 25 30
(2)	2 7 12 17 22 26	(13)	11 7 3 24 20 28	(23)	21 12 3 19 10 30
(3)	3 8 13 18 23 26	(14)	16 12 8 4 25 28	(24)	6 22 13 4 20 30
(4)	4 9 14 19 24 26	(15)	21 17 13 9 5 28	(25)	16 7 23 14 5 30
(5)	5 10 15 20 25 26	(16)	1 12 23 9 20 29	(26)	1 7 13 19 25 31
(6)	1 2 3 4 5 27	(17)	16 2 13 24 10 29	(27)	21 2 8 14 20 31
(7)	6 7 8 9 10 27	(18)	6 17 3 14 25 29	(28)	16 22 3 9 15 31
(8)	11 12 13 14 15 27	(19)	21 7 18 4 15 29	(29)	11 17 23 4 10 31
(9)	16 17 18 19 20 27	(20)	11 22 8 19 5 29	(30)	6 12 18 24 5 31
(10)	21 22 23 24 25 27	(21)	1 17 8 24 15 30	(31)	26 27 28 29 30 31
(11)	1 22 18 14 10 28				

Plan 11.41 $t = 31, k = 10, r = 10, b = 31, \lambda = 3, E = 0.93$, Type IV

Block

(1)	1 6 9 12 17 18 24 26 27 29	(17)	5 6 10 8 18 21 26 28 22 31
(2)	2 7 10 13 18 19 25 27 28 29	(18)	6 7 11 9 19 15 27 22 23 31
(3)	3 1 11 14 19 20 26 28 22 29	(19)	7 1 12 10 20 16 28 23 24 31
(4)	4 2 12 8 20 21 27 22 23 29	(20)	1 2 13 11 21 17 22 24 25 31
(5)	5 3 13 9 21 15 28 23 24 29	(21)	2 3 14 12 15 18 23 25 26 31
(6)	6 4 14 10 15 16 22 24 25 29	(22)	1 2 4 8 9 11 15 16 18 28
(7)	7 5 8 11 16 17 23 25 26 29	(23)	2 3 5 9 10 12 16 17 19 22
(8)	2 5 10 11 15 20 24 26 27 30	(24)	3 4 6 10 11 13 17 18 20 23
(9)	3 6 11 12 16 21 25 27 28 30	(25)	4 5 7 11 12 14 18 19 21 24
(10)	4 7 12 13 17 15 26 28 22 30	(26)	5 6 1 12 13 8 19 20 15 25
(11)	5 1 13 14 18 16 27 22 23 30	(27)	6 7 2 13 14 9 20 21 16 26
(12)	6 2 14 8 19 17 28 23 24 30	(28)	7 1 3 14 8 10 21 15 17 27
(13)	7 3 8 9 20 18 22 24 25 30	(29)	1 2 3 4 5 6 7 29 30 31
(14)	1 4 9 10 21 19 23 25 26 30	(30)	8 9 10 11 12 13 14 29 30 31
(15)	3 4 8 13 16 19 24 26 27 31	(31)	15 16 17 18 19 20 21 29 30 31
(16)	4 5 9 14 17 20 25 27 28 31		

Plan 11.42 $t = 37, k = 9, r = 9, b = 37, \lambda = 2, E = 0.91$, Type IV

Block

(1)	34 13 20 36 3 31 25 33 21	(20)	26 7 16 1 9 33 10 34 12
(2)	12 4 36 15 37 13 10 19 29	(21)	34 20 18 17 15 5 24 4 9
(3)	8 37 32 6 9 15 11 25 33	(22)	30 36 17 29 21 9 27 32 16
(4)	21 2 5 29 4 28 33 11 7	(23)	8 12 20 18 23 21 27 7 37
(5)	36 24 2 34 6 23 28 16 37	(24)	11 16 27 22 12 25 24 4 31
(6)	35 11 18 9 12 36 3 14 28	(25)	32 13 22 7 15 2 16 3 18
(7)	32 12 24 20 35 2 33 30 19	(26)	35 21 19 18 16 6 25 5 10
(8)	2 11 8 34 27 35 10 13 17	(27)	24 29 3 25 37 1 35 7 17
(9)	19 34 29 11 1 18 23 32 31	(28)	12 22 5 8 34 3 6 29 30
(10)	15 21 12 1 2 6 31 17 14	(29)	26 35 21 14 32 4 22 37 34
(11)	17 36 6 22 26 11 7 20 19	(30)	9 14 25 2 10 22 23 20 29
(12)	24 3 26 11 23 10 30 15 21	(31)	19 16 23 17 33 8 3 4 14
(13)	37 16 30 1 13 14 11 20 5	(32)	17 32 28 5 13 12 26 25 23
(14)	3 4 6 20 10 32 27 28 1	(33)	23 36 27 35 15 5 22 1 33
(15)	35 13 9 23 31 30 7 6 4	(34)	36 18 25 4 8 30 26 2 1
(16)	14 27 18 24 6 29 26 13 33	(35)	8 24 10 32 7 31 14 5 36
(17)	8 13 24 9 21 19 28 1 22	(36)	5 26 3 19 27 2 31 37 9
(18)	37 30 31 22 10 28 33 17 18	(37)	28 19 25 30 34 15 7 14 27
(19)	20 29 15 8 26 35 16 28 31		

Plan 11.43 $t = 41, k = 5, r = 10, b = 82, \lambda = 1, E = 0.82$, Type III

Block

(1)	1	37	16	18	10
(2)	2	38	17	19	11
(3)	3	39	18	20	12
(4)	4	40	19	21	13
(5)	5	41	20	22	14
(6)	6	1	21	23	15
(7)	7	2	22	24	16
(8)	8	3	23	25	17
(9)	9	4	24	26	18
(10)	10	5	25	27	19
(11)	11	6	26	28	20
(12)	12	7	27	29	21
(13)	13	8	28	30	22
(14)	14	9	29	31	23
(15)	15	10	30	32	24
(16)	16	11	31	33	25
(17)	17	12	32	34	26
(18)	18	13	33	35	27
(19)	19	14	34	36	28
(20)	20	15	35	37	29
(21)	21	16	36	38	30
(22)	22	17	37	39	31
(23)	23	18	38	40	32
(24)	24	19	39	41	33
(25)	25	20	40	1	34
(26)	26	21	41	2	35
(27)	27	22	1	3	36
(28)	28	23	2	4	37
(29)	29	24	3	5	38
(30)	30	25	4	6	39
(31)	31	26	5	7	40
(32)	32	27	6	8	41
(33)	33	28	7	9	1
(34)	34	29	8	10	2
(35)	35	30	9	11	3
(36)	36	31	10	12	4
(37)	37	32	11	13	5
(38)	38	33	12	14	6
(39)	39	34	13	15	7
(40)	40	35	14	16	8
(41)	41	36	15	17	9

(42)	8	9	5	21	39
(43)	9	10	6	22	40
(44)	10	11	7	23	41
(45)	11	12	8	24	1
(46)	12	13	9	25	2
(47)	13	14	10	26	3
(48)	14	15	11	27	4
(49)	15	16	12	28	5
(50)	16	17	13	29	6
(51)	17	18	14	30	7
(52)	18	19	15	31	8
(53)	19	20	16	32	9
(54)	20	21	17	33	10
(55)	21	22	18	34	11
(56)	22	23	19	35	12
(57)	23	24	20	36	13
(58)	24	25	21	37	14
(59)	25	26	22	38	15
(60)	26	27	23	39	16
(61)	27	28	24	40	17
(62)	28	29	25	41	18
(63)	29	30	26	1	19
(64)	30	31	27	2	20
(65)	31	32	28	3	21
(66)	32	33	29	4	22
(67)	33	34	30	5	23
(68)	34	35	31	6	24
(69)	35	36	32	7	25
(70)	36	37	33	8	26
(71)	37	38	34	9	27
(72)	38	39	35	10	28
(73)	39	40	36	11	29
(74)	40	41	37	12	30
(75)	41	1	38	13	31
(76)	1	2	39	14	32
(77)	2	3	40	15	33
(78)	3	4	41	16	34
(79)	4	5	1	17	35
(80)	5	6	2	18	36
(81)	6	7	3	19	37
(82)	7	8	4	20	38

Block

(1)	1	8	15	22	29	36	43	50
(2)	2	9	16	23	30	37	44	50
(3)	3	10	17	24	31	38	45	50
(4)	4	11	18	25	32	39	46	50
(5)	5	12	19	26	33	40	47	50
(6)	6	13	20	27	34	41	48	50
(7)	7	14	21	28	35	42	49	50
(8)	1	2	3	4	5	6	7	51
(9)	8	9	10	11	12	13	14	51
(10)	15	16	17	18	19	20	21	51
(11)	22	23	24	25	26	27	28	51
(12)	29	30	31	32	33	34	35	51
(13)	36	37	38	39	40	41	42	51
(14)	43	44	45	46	47	48	49	51
(15)	1	44	38	32	26	20	14	52
(16)	8	2	45	39	33	27	21	52
(17)	15	9	3	46	40	34	28	52
(18)	22	16	10	4	47	41	35	52
(19)	29	23	17	11	5	48	42	52
(20)	36	30	24	18	12	6	49	52
(21)	43	37	31	25	19	13	7	52
(22)	1	37	24	11	47	34	21	53
(23)	15	2	38	25	12	48	35	53
(24)	29	16	3	39	26	13	49	53
(25)	43	30	17	4	40	27	14	53
(26)	8	44	31	18	5	41	28	53
(27)	22	9	45	32	19	6	42	53
(28)	36	23	10	46	33	20	7	53
(29)	1	30	10	39	19	48	28	54
(30)	22	2	31	11	40	20	49	54
(31)	43	23	3	32	12	41	21	54
(32)	15	44	24	4	33	13	42	54
(33)	36	16	45	25	5	34	14	54
(34)	8	37	17	46	26	6	35	54
(35)	29	9	38	18	47	27	7	54
(36)	1	23	45	18	40	13	35	55
(37)	29	2	24	46	19	41	14	55
(38)	8	30	3	25	47	20	42	55
(39)	36	9	31	4	26	48	21	55
(40)	15	37	10	32	5	27	49	55
(41)	43	16	38	11	33	6	28	55
(42)	22	44	17	39	12	34	7	55
(43)	1	16	31	46	12	27	42	56
(44)	36	2	17	32	47	13	28	56
(45)	22	37	3	18	33	48	14	56
(46)	8	23	38	4	19	34	49	56
(47)	43	9	24	39	5	20	35	56
(48)	29	44	10	25	40	6	21	56
(49)	15	30	45	11	26	41	7	56
(50)	1	9	17	25	33	41	49	57
(51)	43	2	10	18	26	34	42	57
(52)	36	44	3	11	19	27	35	57
(53)	29	37	45	4	12	20	28	57
(54)	22	30	38	46	5	13	21	57
(55)	15	23	31	39	47	6	14	57
(56)	8	16	24	32	40	48	7	57
(57)	50	51	52	53	54	55	56	57

Plan 11.45 $t = 73, k = 9, r = 9, b = 73, \lambda = 1, E = 0.90$, Type IV

Block

(1)	1 9 17 25 33 41 49 57 65	(38)	49 58 35 28 13 6 23 48 69
(2)	2 10 18 26 34 42 50 58 65	(39)	9 42 27 36 53 22 7 64 69
(3)	3 11 19 27 35 43 51 59 65	(40)	33 18 51 12 29 46 63 8 69
(4)	4 12 20 28 36 44 52 60 65	(41)	1 34 11 20 53 30 63 48 70
(5)	5 13 21 29 37 45 53 61 65	(42)	33 2 59 28 45 22 15 56 70
(6)	6 14 22 30 38 46 54 62 65	(43)	9 58 3 52 21 46 39 32 70
(7)	7 15 23 31 39 47 55 63 65	(44)	17 26 51 4 13 38 47 64 70
(8)	8 16 24 32 40 48 56 64 65	(45)	49 42 19 12 5 62 31 40 70
(9)	1 2 3 4 5 6 7 8 66	(46)	25 18 43 36 61 6 55 16 70
(10)	9 10 11 12 13 14 15 16 66	(47)	57 10 35 44 29 54 7 24 70
(11)	17 18 19 20 21 22 23 24 66	(48)	41 50 27 60 37 14 23 8 70
(12)	25 26 27 28 29 30 31 32 66	(49)	1 42 59 52 29 38 23 16 71
(13)	33 34 35 36 37 38 39 40 66	(50)	41 2 19 36 13 54 63 32 71
(14)	41 42 43 44 45 46 47 48 66	(51)	57 18 3 28 53 14 47 40 71
(15)	49 50 51 52 53 54 55 56 66	(52)	49 34 27 4 61 46 15 24 71
(16)	57 58 59 60 61 62 63 64 66	(53)	25 10 51 60 5 22 39 48 71
(17)	1 10 19 28 37 46 55 64 67	(54)	33 50 11 44 21 6 31 64 71
(18)	9 2 51 44 61 30 23 40 67	(55)	17 58 43 12 37 30 7 56 71
(19)	17 50 3 36 29 62 15 48 67	(56)	9 26 35 20 45 62 55 8 71
(20)	25 42 35 4 21 14 63 56 67	(57)	1 50 35 12 61 22 47 32 72
(21)	33 58 27 20 5 54 47 16 67	(58)	49 2 43 20 29 14 39 64 72
(22)	41 26 59 12 53 6 39 24 67	(59)	33 42 3 60 13 30 55 24 72
(23)	49 18 11 60 45 38 7 32 67	(60)	9 18 59 4 37 54 31 48 72
(24)	57 34 43 52 13 22 31 8 67	(61)	57 26 11 36 5 46 23 56 72
(25)	1 18 27 44 13 62 39 56 68	(62)	17 10 27 52 45 6 63 40 72
(26)	17 2 35 60 53 46 31 16 68	(63)	41 34 51 28 21 62 7 16 72
(27)	25 34 3 12 45 54 23 64 68	(64)	25 58 19 44 53 38 15 8 72
(28)	41 58 11 4 29 22 55 40 68	(65)	1 58 51 36 45 14 31 24 73
(29)	9 50 43 28 5 38 63 24 68	(66)	57 2 27 12 21 38 55 48 73
(30)	57 42 51 20 37 6 15 32 68	(67)	49 26 3 44 37 22 63 16 73
(31)	33 26 19 52 61 14 7 48 68	(68)	33 10 43 4 53 62 23 32 73
(32)	49 10 59 36 21 30 47 8 68	(69)	41 18 35 52 5 30 15 64 73
(33)	1 26 43 60 21 54 15 40 69	(70)	9 34 19 60 29 6 47 56 73
(34)	25 2 11 52 37 62 47 24 69	(71)	25 50 59 20 13 46 7 40 73
(35)	41 10 3 20 61 38 31 56 69	(72)	17 42 11 28 61 54 39 8 73
(36)	57 50 19 4 45 30 39 16 69	(73)	65 66 67 68 69 70 71 72 73
(37)	17 34 59 44 5 14 55 32 69		

Plan 11.46 $t = 91, k = 10, r = 10, b = 91, \lambda = 1, E = 0.91$, Type IV

Block

(1)	1 10 19 28 37 46 55 64 73 82	(47)	28 2 57 40 14 69 52 26 81 87
(2)	2 11 20 29 38 47 56 65 74 82	(48)	55 29 3 67 41 15 79 53 27 87
(3)	3 12 21 30 39 48 57 66 75 82	(49)	19 74 48 4 59 33 16 71 45 87
(4)	4 13 22 31 40 49 58 67 76 82	(50)	46 20 75 31 5 60 43 17 72 87
(5)	5 14 23 32 41 50 59 68 77 82	(51)	73 47 21 58 32 6 70 44 18 87
(6)	6 15 24 33 42 51 60 69 78 82	(52)	10 65 39 22 77 51 7 62 36 87
(7)	7 16 25 34 43 52 61 70 79 82	(53)	37 11 66 49 23 78 34 8 63 87
(8)	8 17 26 35 44 53 62 71 80 82	(54)	64 38 12 76 50 24 61 35 9 87
(9)	9 18 27 36 45 54 63 72 81 82	(55)	1 47 66 76 14 33 43 62 27 88
(10)	1 2 3 4 5 6 7 8 9 83	(56)	64 2 48 31 77 15 25 44 63 88
(11)	10 11 12 13 14 15 16 17 18 83	(57)	46 65 3 13 32 78 61 26 45 88
(12)	19 20 21 22 23 24 25 26 27 83	(58)	37 56 21 4 50 69 79 17 36 88
(13)	28 29 30 31 32 33 34 35 36 83	(59)	19 38 57 67 5 51 34 80 18 88
(14)	37 38 39 40 41 42 43 44 45 83	(60)	55 20 39 49 68 6 16 35 81 88
(15)	46 47 48 49 50 51 52 53 54 83	(61)	73 11 30 40 59 24 7 53 72 88
(16)	55 56 57 58 59 60 61 62 63 83	(62)	28 74 12 22 41 60 70 8 54 88
(17)	64 65 66 67 68 69 70 71 72 83	(63)	10 29 75 58 23 42 52 71 9 88
(18)	73 74 75 76 77 78 79 80 81 83	(64)	1 74 39 67 32 24 52 17 63 89
(19)	1 20 12 58 77 69 34 53 45 84	(65)	37 2 75 22 68 33 61 53 18 89
(20)	10 2 21 67 59 78 43 35 54 84	(66)	73 38 3 31 23 69 16 62 54 89
(21)	19 11 3 76 68 60 52 44 36 84	(67)	46 11 57 4 77 42 70 35 27 89
(22)	28 47 39 4 23 15 61 80 72 84	(68)	55 47 12 40 5 78 25 71 36 89
(23)	37 29 48 13 5 24 70 62 81 84	(69)	10 56 48 76 41 6 34 26 72 89
(24)	46 38 30 22 14 6 79 71 63 84	(70)	64 29 21 49 14 60 7 80 45 89
(25)	55 74 66 31 50 42 7 26 18 84	(71)	19 65 30 58 50 15 43 8 81 89
(26)	64 56 75 40 32 51 16 8 27 84	(72)	28 20 66 13 59 51 79 44 9 89
(27)	73 65 57 49 41 33 25 17 9 84	(73)	1 65 48 40 23 60 79 35 18 90
(28)	1 11 21 31 41 51 61 71 81 85	(74)	46 2 66 58 41 24 16 80 36 90
(29)	19 2 12 49 32 42 79 62 72 85	(75)	64 47 3 22 59 42 34 17 81 90
(30)	10 20 3 40 50 33 70 80 63 85	(76)	73 29 12 4 68 51 43 26 63 90
(31)	55 65 75 4 14 24 34 44 54 85	(77)	10 74 30 49 5 69 61 44 27 90
(32)	73 56 66 22 5 15 52 35 45 85	(78)	28 11 75 67 50 6 25 62 45 90
(33)	64 74 57 13 23 6 43 53 36 85	(79)	37 20 57 76 32 15 7 71 54 90
(34)	28 38 48 58 68 78 7 17 27 85	(80)	55 38 21 13 77 33 52 8 72 90
(35)	46 29 39 76 59 69 25 8 18 85	(81)	19 56 39 31 14 78 70 53 9 90
(36)	37 47 30 67 77 60 16 26 9 85	(82)	1 38 75 49 59 15 70 26 36 91
(37)	1 29 57 22 50 78 16 44 72 86	(83)	73 2 39 13 50 60 34 71 27 91
(38)	55 2 30 76 23 51 70 17 45 86	(84)	37 74 3 58 14 51 25 35 72 91
(39)	28 56 3 49 77 24 43 71 18 86	(85)	64 20 30 4 41 78 52 62 18 91
(40)	10 38 66 4 32 60 25 53 81 86	(86)	28 65 21 76 5 42 16 53 63 91
(41)	64 11 39 58 5 33 79 26 54 86	(87)	19 29 66 40 77 6 61 17 54 91
(42)	37 65 12 31 59 6 52 80 27 86	(88)	46 56 12 67 23 33 7 44 81 91
(43)	19 47 75 13 41 69 7 35 63 86	(89)	10 47 57 31 68 24 79 8 45 91
(44)	73 20 48 67 14 42 61 8 36 86	(90)	55 11 48 22 32 69 43 80 9 91
(45)	46 74 21 40 68 15 34 62 9 86	(91)	82 83 84 85 86 87 88 89 90 91
(46)	1 56 30 13 68 42 25 80 54 87		

CHAPTER 12

LATTICE SQUARES

12.1 Description

12.11 Balanced Lattice Squares. The number of treatments must be an exact square. Within each replicate, the k^2 treatments are arranged on the plan in a $k \times k$ square. The method of grouping into rows and columns, which varies in successive replications, is such that the treatment means can be adjusted for differences among the rows and columns of each square. Thus, in addition to the elimination of differences among replicates from the experimental errors, the design permits a "double control" within each replicate, similar to that obtained in a latin square.

The principle which governs the grouping into rows and columns may be seen from the plans 12.1-12.8. With 9 treatments, for instance, treatment 1 occurs in the same row as treatments 2, 3, 6, and 8, and in the same column as treatments 4, 5, 7, and 9. Thus every other treatment appears with treatment 1 either in the same row or in the same column. More generally, any pair of treatments occurs once in the same row or in the same column.

This property holds for all plans having an odd number of treatments, i.e., for 9, 25, 49, 81, 121, and 169 treatments. The number of replicates, for k^2 treatments, is $(k + 1)/2$. The standard error of the difference is not exactly the same for all pairs of treatments, though the variation in accuracy is small. Separate formulae are given for the two standard errors for cases in which they may be needed.

A design with twice the basic number of replicates (e.g., 25 treatments in 6 replicates) is obtained by repeating the plan, *the rows being interchanged with the columns*. In designs so produced, every pair of treatments occurs together once in the same row *and* once in the same column. On account of this property, the standard error of the difference is the same for all pairs of treatments, whether the row and column differences are large or small.

Within the most useful range, the only even numbers of treatments which provide lattice square designs are 16 and 64. These designs have, respectively, 5 and 9, i.e. $(k + 1)$ replicates. Every pair of treatments occurs once both in the same row and in the same column.

The available selection of designs (up to 12 replicates) is summarized

in table 12.1. By analogy with balanced lattices, the designs may be called *balanced* lattice squares.

TABLE 12.1 AVAILABLE NUMBERS OF TREATMENTS AND REPLICATIONS FOR
BALANCED LATTICE SQUARES

Number of treatments	9	16	25	49	64	81	121	169
Number of replicates	4, 8	5, 10	3, 6	4, 8	9	5, 10	6, 12	7

For 16 treatments in 10 replicates, the basic plan is used twice. Although the basic plan (plan 12.1) requires only 2 replicates, the design for 9 treatments is not recommended with less than 4 replicates. Even in this case there are only 8 d.f. for estimating row and column variances and only 8 d.f. for error. Little would be gained over randomized blocks unless the extra controls were highly effective. A design with 9 treatments and 8 replicates is found by repeating the design for 9 treatments and 4 replicates.

Certain factorial experiments can be arranged in *balanced* lattice squares; e.g., 3×3 , 8×2 , 4×4 , $4 \times 2 \times 2$, 2^4 , 5×5 , 7×7 , 8×8 , 4^3 , 2^6 , 9×9 , 3^4 . All main effects and interactions are confounded to the same extent with rows and columns.

12.12 Partially Balanced Squares. For a balanced design, the available numbers of replications are rather severely restricted. Although designs are possible for other numbers of replicates, they lack the symmetry of the designs described above, with the consequence that the statistical analysis usually becomes more complicated. It happens, however, that if the number of replications is less than that required for a balanced design, the statistical analysis follows the same procedure as for balanced designs, apart from minor changes in some formulae. In experimentation where double grouping has proved effective but where the replications are not sufficient for a balanced lattice square, the additional designs shown in table 12.2 may be useful.

TABLE 12.2 OTHER LATTICE SQUARE DESIGNS

Number of treatments	49	64	81	121	169
Number of replicates	3	3, 4	3, 4	3, 4, 5	3, 4, 5, 6

When the number of treatments is odd, the plans for these designs are obtained simply by taking the desired number of replicates from the plans for the balanced designs. Thus, for 49 treatments in threefold replication, we use squares I to III from plan 12.4. With 64 treatments, squares I, III, and V are used for 3 replicates, and squares I, III, V, and VII for 4 replicates.

Partially balanced lattice squares are analogous to the triple, quadruple, and quintuple lattices.

12.13 Arrangement of Experimental Material. The method of arranging the experimental material is similar to that for an ordinary latin square. With k^2 treatments, the units are arranged in $k \times k$ squares so that the rows and columns of each square correspond to the two types of variation whose effects we wish to eliminate from the errors. In field experiments, the plots of each replicate are usually laid out in square formation, in which case row and column differences represent fertility variations in two directions at right angles to each other. If the width of a greenhouse bench accommodates 5 pots, 25 treatments can be laid out in replicates which consist of 5 rows of 5 pots each.

12.14 Randomization. Within each replicate the rows and columns of the basic plan should be permuted separately at random before applying the treatments. It is also advisable to assign the treatments at random to the treatment numbers in the plan.

12.2 Statistical Analysis

12.21 Designs with $(k + 1)/2$, or Fewer, Replications. The same instructions apply to the partially balanced designs and to the balanced designs with $(k + 1)/2$ replications. Table 12.3 shows the plan and yields for a lattice square with 25 corn hybrids in 3 replications, conducted by the North Carolina Experiment Station in 1942. Each plot contained 4 rows, with 10 hills per row.

The computations proceed as follows. The number of treatments is k^2 , and the number of replications r .

1. Find the row, column, treatment, replication, and grand totals. The treatment totals are placed in a square under the plan.

2. For each row, calculate the L value, where

$$L = \text{total (from all replications) of all treatments included in the row} - r \text{ times row total}$$

e.g.,

$$L_1 = 89.8 + 93.3 + 82.1 + 88.9 + 91.1 - 3(159.1) = -32.1$$

As a check, the total of the L values over a replication equals

$$(\text{Grand total}) - r(\text{replication total})$$

Thus,

$$-76.2 = 2189.4 - 3(755.2)$$

Similarly for each column we obtain M values.

$$M = \text{total (from all replications) of all treatments included in the column} - r \text{ times column total}$$

TABLE 12.3 YIELDS (POUNDS CORN PER PLOT) FOR A 5 X 5 LATTICE SQUARE IN 3 REPLICATIONS

Treatment numbers are shown in parentheses										Total	L	s
Square I												
(18) 33.3	(9) 30.7	(11) 35.4	(2) 30.1	(25) 29.6						159.1	-32.1	-2.58
(24) 24.6	(15) 30.8	(17) 28.8	(8) 34.8	(1) 32.5						151.5	-10.5	-0.84
(12) 28.5	(3) 24.0	(10) 28.4	(21) 25.0	(19) 35.1						141.0	+22.4	+1.80
(6) 26.7	(22) 27.2	(4) 25.6	(20) 25.0	(13) 29.4						133.9	+11.0	+0.88
(5) 40.1	(16) 35.7	(23) 30.1	(14) 30.3	(7) 33.5						169.7	-67.0	-5.39
Total	153.2	148.4	148.3	145.2	160.1					755.2		-6.13
M	-16.0	+3.8	-12.8	-10.0	-41.2						-76.2	
e	- 0.83	+0.20	- 0.66	- 0.52	- 2.13					-3.94		
Square II										Total	L	s
(20) 30.9	(17) 33.3	(19) 38.8	(16) 27.7	(18) 34.4								
(15) 37.2	(12) 31.2	(14) 27.9	(11) 27.3	(13) 21.6						145.2	- 5.8	- 0.47
(25) 32.7	(22) 43.0	(24) 28.5	(21) 24.7	(23) 22.7						151.6	- 46.2	- 3.71
(5) 32.0	(2) 32.8	(4) 31.8	(1) 28.7	(3) 32.3						157.6	- 33.8	- 2.72
(10) 39.8	(7) 37.3	(9) 31.9	(6) 34.0	(8) 34.3						177.3	- 69.3	- 5.57
Total	172.6	177.6	158.9	142.4	145.3					796.8		-16.16
M	-62.2	-85.5	-30.1	-12.6	-10.6						-201.0	
e	- 3.22	- 4.43	- 1.56	- 0.65	- 0.55					-10.41		
Square III										Total	L	s
(19) 28.7	(15) 26.3	(23) 21.7	(6) 21.9	(2) 26.0								
(11) 19.4	(7) 17.3	(20) 16.9	(3) 22.6	(24) 24.2						100.4	+ 98.0	+ 7.88
(22) 18.3	(18) 22.1	(1) 17.5	(14) 25.0	(10) 26.9						109.8	+105.9	+ 8.51
(5) 30.2	(21) 27.5	(9) 30.7	(17) 28.1	(13) 27.6						144.1	+ 9.3	+ 0.75
(8) 34.4	(4) 32.8	(12) 31.9	(25) 28.8	(16) 30.6						158.5	- 5.1	- 0.41
Total	131.0	126.0	118.7	126.4	135.3					637.4		+22.23
M	+86.0	+61.6	+54.8	+46.8	+28.0						+277.2	
e	+ 4.45	+ 3.19	+ 2.84	+ 2.42	+ 1.45					+14.35		
Grand total										2189.4	0	0
Treatment totals												
(1) 78.7	(2) 88.9	(3) 78.9	(4) 90.2	(5) 102.3								
(6) 82.6	(7) 88.1	(8) 103.5	(9) 93.3	(10) 95.1								
(11) 82.1	(12) 91.6	(13) 78.6	(14) 83.2	(15) 94.3								
(16) 94.0	(17) 90.2	(18) 89.8	(19) 102.6	(20) 72.8								
(21) 77.2	(22) 88.5	(23) 74.5	(24) 77.3	(25) 91.1						2189.4		
Adjusted treatment totals												
(1) 83.7	(2) 85.7	(3) 87.9	(4) 88.9	(5) 95.3								
(6) 84.4	(7) 81.6	(8) 100.1	(9) 87.4	(10) 97.4								
(11) 90.1	(12) 90.1	(13) 78.5	(14) 86.2	(15) 98.7								
(16) 85.5	(17) 83.8	(18) 93.8	(19) 107.0	(20) 77.0								
(21) 78.1	(22) 94.4	(23) 72.6	(24) 79.7	(25) 81.5								

3. We now compute the analysis of variance. The total s.s. and the sums of squares for replications and treatments are found in the usual way. If the symbol L_r denotes a replication total of the L 's, the sum of squares for rows within replications, adjusted for treatments, is

$$\begin{aligned} & \frac{\sum L^2}{kr(r-1)} - \frac{\sum L_r^2}{k^2r(r-1)} \\ &= \frac{(32.1)^2 + (10.5)^2 + \cdots + (5.1)^2}{30} - \frac{(76.2)^2 + (201.0)^2 + (277.2)^2}{150} \\ &= 1405.95 - 820.32 = 585.63 \end{aligned}$$

The sum of squares for columns, eliminating treatments, is

$$\begin{aligned} & \frac{(16.0)^2 + (3.8)^2 + \cdots + (28.0)^2}{30} - \frac{(76.2)^2 + (201.0)^2 + (277.2)^2}{150} \\ &= 1058.53 - 820.32 = 238.21 \end{aligned}$$

The complete analysis of variance is shown in table 12.4.

TABLE 12.4 ANALYSIS OF VARIANCE FOR A 5×5 LATTICE SQUARE

	d.f.		s.s.	m.s.
Replications	$(r-1)$	2	546.88	273.44
Treatments	(k^2-1)	24	611.09	25.46
Rows (adj.)	$r(k-1)$	12	585.63	48.80 E_r
Columns (adj.)	$r(k-1)$	12	238.21	19.85 E_c
Error	$(k-1)(rk-r-k-1)$	24	229.79	9.57 E_e
Total	(rk^2-1)	74	2211.60	

4. This step leads to the adjusted treatment means. Let

E_r = mean square for rows

E_c = mean square for columns

E_e = mean square for error

$$\lambda' = \frac{(E_r - E_e)}{k(r-1)E_r} = \frac{48.80 - 9.57}{(5)(2)(48.80)} = 0.0804$$

$$\mu' = \frac{(E_c - E_e)}{k(r-1)E_c} = \frac{19.85 - 9.57}{(5)(2)(19.85)} = 0.0518$$

If E_r or E_c is less than E_e , then λ' or μ' , as the case may be, is taken as 0.

5. Multiply the L 's by λ' to give δ values, and the M 's by μ' to give ϵ values, as presented in table 12.3.

6. The adjusted total for any treatment is secured by adding to the unadjusted total the δ and ϵ values for each row and column in which the treatment appears. For treatment (1), we have

$$78.7 + (-0.84) + (-2.13) + (-2.72) + (-0.65) + (8.51) + (2.84) = 83.7$$

To obtain the treatment means (not shown) we divide by the number of replicates.

7. The *average variance* of the difference between two adjusted treatment means is

$$\frac{2E_e}{r} \left[1 + \frac{rk}{(k+1)} (\lambda' + \mu') \right] = \frac{2(9.57)}{3} \left[1 + \frac{(3)(5)}{6} (0.1322) \right] \\ = 8.49; \text{ s.e.} = 2.91$$

Except perhaps with the 5×5 and smaller squares, it is sufficiently accurate to use this figure for comparisons between any pair of treatments. More accurately, for two treatments that appear in the same row, the variance of the difference is

$$\frac{2E_e}{r} [1 + (r-1)\lambda' + r\mu'] = 8.40; \text{ s.e.} = 2.90$$

For two treatments in the same column.

$$\frac{2E_e}{r} [1 + r\lambda' + (r-1)\mu'] = 8.58; \text{ s.e.} = 2.93$$

Clearly the average error is good enough in this experiment. For the partially balanced designs one more formula is needed, since some pairs of treatments do not occur together either in a row or in a column. The variance of the difference is

$$\frac{2E_e}{r} [1 + r(\lambda' + \mu')]$$

8. As is typical of these designs, the analysis does not provide an exact F -test. The treatments m.s. in the analysis (table 12.4) cannot be compared with the error m.s., since the former has not been adjusted for row and column effects. For an approximate F -test, compute the sum of squares of deviations of the adjusted treatment totals in table 12.3. This comes to 1606.05. Division by 3, since there are 3 replicates, gives

535.35, with a mean square of 22.31. This may be compared with the effective error m.s., which is

$$E_e \left[1 + \frac{rk}{k+1} (\lambda' + \mu') \right] = 9.57 \left[1 + \frac{15}{6} (0.1322) \right] = 12.73$$

The F -ratio is 22.31/12.73, or 1.75, with 24 and 24 d.f.

9. The gain in precision relative to randomized blocks is estimated as follows. If the experiment were analyzed as randomized blocks, rows and columns would be amalgamated with error, giving

	d.f.	s.s.	m.s.
Replications	2	546.88	
Treatments	24	611.09	
Error	48	1053.63	21.95

This error, 21.95, is compared with the effective error m.s., 12.73. The relative information is estimated as 21.95/12.73, or 172%. Thus 3 replicates of the lattice square appear about as precise as 5 with randomized blocks.

Often a comparison with a lattice design will be of more interest. This experiment could have been planned as a triple lattice, and in that event the rows would probably have been chosen as blocks. The intra-block error would be derived from the pooled sum of squares for columns and error in table 12.4. The mean square would be 13.00, with 36 d.f. Consequently, for the triple lattice we have $E_b = 48.80$ (m.s. for rows), $E_e = 13.00$. The effective error m.s. (section 10.28) is

$$\begin{aligned} E_e \left[1 + \frac{rk\mu}{k+1} \right] &= E_e \left[1 + \frac{r(E_b - E_e)}{(k+1)(r-1)E_b} \right] \\ &= 13.00 \left[1 + \frac{3(48.80 - 13.00)}{(6)(2)(48.80)} \right] = 15.38 \end{aligned}$$

The relative accuracy of the lattice square to the triple lattice is estimated as 15.38/12.73, or 121%.

An account of the theory and a worked example are given by Yates (12.1) for balanced squares and by Cochran (12.2) for partially balanced squares.

12.22 Designs with $(k+1)$ Replications. In this case the analysis is slightly different, since it is possible to apply a single adjustment for all the rows in which a treatment lies, instead of making a separate adjust-

ment for every row. On the whole, however, the analysis is more complex than with $(k + 1)/2$ replications.

The example is a 4×4 lattice square conducted by the U. S. Bureau of Entomology and Plant Quarantine at Florence, S. C., and described by Wadley (12.3). The treatments were 16 arsenical insecticides applied to cotton with a hand dusting machine. Plots were 10 rows wide and 70 feet long, being about $\frac{1}{18}$ acre. To allow for border effects, records were taken only from the 4 center rows. The data in table 12.5 are the percentages of squares * showing attack by boll weevils. These figures were obtained by examining 100 squares per plot, 25 from each of the 4 center rows. Such counts were made at intervals during the summer: the data are averages from 3 counts made in August.

The simplest computational routine, devised by Yates (12.1), will be presented here, though it is not too well adapted for making clear the meaning of the various steps. Yates's paper should be consulted for an account of the theory.

TABLE 12.6 ORIGINAL AND ADJUSTED TREATMENT TOTALS

	Treat- ment totals, <i>T</i>	<i>R_i</i>	<i>C_i</i>	<i>D</i>	<i>L'</i>	<i>J</i>	<i>K</i>	<i>M'</i>	Adj. treat- ment totals	Adj. means
1	24.6	181.3	164.1	17.2	64.3	81.5	133.1	150.3	32.24	6.45
2	74.7	284.4	196.6	87.8	-250.8	-163.0	100.4	188.2	68.41	13.68
3	33.6	146.4	221.9	-75.5	274.8	199.3	-27.2	-102.7	43.64	8.73
4	56.0	214.2	223.1	-7.9	25.4	17.5	-6.2	-14.1	56.79	11.36
5	50.0	220.7	223.1	-2.4	-31.1	-33.5	-40.7	-43.1	47.20	9.44
6	34.0	211.0	161.4	49.6	-46.6	3.0	151.8	201.4	37.39	7.53
7	33.3	180.0	211.0	-31.0	105.6	74.6	-18.4	-49.4	36.85	7.37
8	48.3	213.4	224.0	-10.6	-1.4	-12.0	-43.8	-54.4	46.58	9.32
9	50.0	197.8	240.3	-42.5	83.4	40.9	-86.6	-129.1	50.07	10.01
10	81.4	259.3	253.5	5.8	-98.5	-92.7	-75.3	-69.5	74.57	14.91
11	98.0	295.1	252.5	42.6	-211.1	-168.5	-40.7	1.9	87.95	17.59
12	68.3	250.1	227.6	22.5	-104.9	-82.4	-14.9	7.6	63.51	12.70
13	60.8	236.0	251.2	-15.2	-64.4	-79.6	-125.2	-140.4	53.45	10.69
14	74.4	265.5	204.4	61.1	-157.5	-96.4	86.9	148.0	71.36	14.27
15	43.3	178.7	236.5	-57.8	152.1	94.3	-79.1	-136.9	46.42	9.28
16	41.7	155.7	199.4	-43.7	260.7	217.0	85.9	42.2	55.46	11.09
Totals	872.4	3489.6	3489.6	0	0	0	0	0	872.39	174.47

1. Find the row, column, replication, and grand totals, all shown in table 12.5.

2. Find the treatment totals *T* (table 12.6). For each treatment find

R_i = total of all rows in which the treatment appears

C_i = total of all columns in which the treatment appears

* A "square" is the name given to the young flower bud.

It may save time to find all 3 quantities, T , R_t , and C_t , by simultaneous addition, since this means that the positions of a treatment in the different replicates are found only once. This can be done if the machine carriage is moved so as to accommodate 3 sets of running totals. The sum of the R_t values and the sum of the C_t values should each equal k times the sum of the T values.

Thereafter the successive columns in table 12.6 are filled out as follows.

$$D = R_t - C_t$$

$$L' = kT - (k + 1)R_t + G$$

$$J = D + L'$$

$$K = J + (k - 1)D$$

$$M' = D + K$$

Note that the total of each of these quantities is zero.

3. Compute the analysis of variance. The only sums of squares requiring special instructions are those for rows and columns. After adjustment for treatment effects, it happens unfortunately that the sums of squares for rows and columns are not mutually orthogonal. By a well-known result in the analysis of variance, their combined sum of squares may be expressed in either of two ways:

$$\begin{aligned} &\text{Rows (adj. for treatments) + columns (adj. for treatments and rows)} \\ &= \text{columns (adj. for treatments) + rows (adj. for treatments and columns)} \end{aligned}$$

Both expressions are computed.

Sum of squares for rows adjusted for treatments

$$= \frac{S(L')^2}{k^3(k + 1)} = \frac{(64.3)^2 + (250.8)^2 + \cdots + (260.7)^2}{320} = 1093.02$$

Sum of squares for rows adjusted for treatments and columns

$$= \frac{S(J^2)}{k^3(k - 1)} = \frac{(81.5)^2 + (163.0)^2 + \cdots + (217.0)^2}{192} = 1026.76$$

$$\text{Sum of squares for columns adjusted for treatments} = \frac{S(M')^2}{k^3(k + 1)} = 625.85$$

Sum of squares for columns adjusted for treatments and rows

$$= \frac{S(K^2)}{k^3(k - 1)} = 559.59$$

Note that $1093.02 + 559.59 = 1026.76 + 625.85$.

TABLE 12.7. ANALYSIS OF VARIANCE

	d.f.		s.s.	m.s.
Replications	k	4	31.56	7.89
Treatments	$k^2 - 1$	15	1244.20	82.95
Rows (adj. for treatments)	$k^2 - 1$	15	1093.02	
Rows (adj. for tr. and col.)	$k^2 - 1$	15	1026.76	$68.45E_r$
Columns (adj. for tr.)	$k^2 - 1$	15	625.85	
Columns (adj. for tr. and rows)	$k^2 - 1$	15	559.59	$37.31E_c$
Error	$(k^2 - 1)(k - 2)$	30	680.17	$22.67E_e$
Total	$k^3 + k^2 - 1$	79	3608.54	

In finding the error s.s. by subtraction, note that we subtract the combined effect of rows and columns *only once*.

4. This leads to the adjusted treatment totals. Compute

$$q = k^2 E_r E_c - E_e^2 = (16)(68.45)(37.31) - (22.67)^2 = 40,348$$

$$Q = (k - 1)q = 121,044$$

$$\lambda' = \frac{(E_r - E_e)(kE_c - E_e)}{Q} = \frac{(45.78)(126.57)}{121,044} = 0.04787$$

$$\mu' = \frac{(E_c - E_e)(kE_r - E_e)}{Q} = \frac{(14.64)(251.13)}{121,044} = 0.03037$$

The adjusted treatment totals are

$$T + \lambda' L' + \mu' M'$$

and are inserted on the right of the M' column in table 12.6. For the means, divide by r .

5. The error variance of the difference between two adjusted means is

$$\frac{2E_e}{r} [1 + k(\lambda' + \mu')] = \frac{2(22.67)}{5} [1 + 4(0.07824)] = 11.9$$

To obtain an approximate F -test of the treatments, compute the sum of squares of deviations of the adjusted treatment totals, and divide by $(k + 1)(k^2 - 1)$ to give the mean square on a single unit basis. This is tested against the effective error m.s., $E_e[1 + k(\lambda' + \mu')]$.

The gain in precision relative to a randomized blocks or lattice design is estimated by the procedure for the lattice square with $(k + 1)/2$ replications.

12.23 Designs with $2(k+1)$ Replications. This case is unlikely to be of much practical interest except possibly with the 3×3 design in 8 replicates and the 4×4 in 10. The analysis is similar to that for $(k+1)$ replications; the changes will be noted very briefly.

The partition of degrees of freedom is as follows.

	d.f.
Replications	$(2k+1)$
Treatments	(k^2-1)
Rows	
Component (a)	(k^2-1)
Component (b)	(k^2-1)
Columns	
Component (a)	(k^2-1)
Component (b)	(k^2-1)
Error	$(2k-3)(k^2-1)$
Total	$2k^2(k+1)-1$

Corresponding to any row, there is another row with the same set of treatments. Find the differences D_r between the totals of corresponding rows. The sum of squares of deviations of the D_r from their replication means, divided by $2k$, gives component (a) of the rows s.s. Component (b) corresponds to the component that was obtained with $(k+1)$ replicates. As before, it has two forms, an L' and a J form. Both are found in the same way as with $(k+1)$ replications, except that the divisors must be doubled. The sums of squares for columns are derived similarly.

The formulae for the adjustment factors are a little more complicated. Let E_r be the mean square found by pooling component (a) and the J component of the rows s.s., with an analogous definition for E_c . Calculate

$$W_i = \frac{1}{E_c}; \quad W_r = \frac{(2k-1)}{2kE_r - E_c}; \quad W_c = \frac{(2k-1)}{2kE_c - E_e}$$

$$\lambda' = \frac{(W_i - W_r)}{k[W_r + W_c + (k-1)W_i]}; \quad \mu' = \frac{W_i - W_c}{k[W_r + W_c - (k-1)W_i]}$$

The remainder of the analysis proceeds without change.

12.24 Missing Data. The procedure for analyzing lattice squares when certain observations are missing has been worked out by Cornish (12.4). As in the case of lattices (section 10.13), the method requires two different estimates of each missing observation, one being that found by minimizing the intra-row-and-column error, the other being that appropriate to the case where the data are analyzed as a randomized block

experiment. In order to reduce the amount of computation, we present a cruder method which should be satisfactory if only a small fraction of the total observations is missing.

In this approach we estimate each missing observation by the first method mentioned above, that is, by minimizing the intra-row-and-column error s.s. Thereafter the analysis proceeds as usual: 1 d.f. is deleted from the total and intra-row-and-column error s.s. for each missing value. Special rules for *t*-tests are given later.

The following totals are used in the formulae for inserting estimates in place of the missing values. *R*, *C*, *T*, and *P* are the totals of the row, column, treatment, and replication, respectively, that contain the missing observation, while *G* is the grand total. Further,

S_x = total of all *other* rows and columns in which the treatment with the missing value appears (note that this sum does not include the row and column which contain the missing value)

T_x = total (from all replicates) of all *other* treatments which appear in the row or column that contains the missing value (this total does not include the treatment that has the missing value)

Since it is important that the definitions of S_x and T_x be correctly understood, it is worth working through the numerical example given below.

Experiment with $(k + 1)/2$ replications

$$x = \frac{k(k-1)(R+C) - (k+3)P + 2k(k-2)T + 6G - 2kS_x - 2kT_x}{(k-1)^2(k-3)}$$

Experiment with $(k + 1)$ replications

$$x = \frac{k(k-1)(R+C) - (k+1)P + k(k-1)T + 3G - kS_x - kT_x}{(k-1)^2(k-2)}$$

Experiment with less than $(k + 1)/2$ replications. Because of the lack of symmetry in this design, an additional symbol is needed.

U_x = total (from all replicates) of all treatments that appear in the same row or column as the treatment that has the missing value.

The distinction between U_x and T_x should be realized. T_x is a total over only those other treatments that are in the actual row or column that contains the missing value, while U_x is a total over all other treatments that appear somewhere in the experiment in the same row or column as the treatment that has the missing value. Thus T_x is a part of U_x . It may help to note that T_x is a total over $2(k-1)$ treatments,

or $2r(k-1)$ observations, where r is the number of replications, while U_x is a total over $2r(k-1)$ treatments, or $2r^2(k-1)$ observations.

$$x = \frac{\left\{ \begin{aligned} &kr(r-1)(R+C) - r(r+1)P + k(k-2)(r-1)T \\ &+ (r+1)G - krS_x - krT_x + kU_x \end{aligned} \right\}}{(k-1)(r-1)(kr-k-r-1)}$$

Example. In the corn experiment in table 12.3, suppose that treatment (18) in replication I is missing. Here $k = 5$, $r = (k+1)/2 = 3$. Omitting the observation 33.3 for this treatment, we find

$$R = 125.8; \quad C = 119.9; \quad R + C = 245.7; \quad P = 721.9$$

$$T = 56.5; \quad G = 2156.1$$

S_x is the total of the rows and columns in which treatment (18) appears in squares II and III. Thus

$$S_x = 165.1 + 145.3 + 109.8 + 126.0 = 546.2$$

Finally, T_x is the total for treatments (2), (9), (11), and (25), which appear in the row that has the missing value, plus that for treatments (5), (6), (12), and (24) which appear in the column with the missing value. These totals are to be taken from the table of treatment totals. This gives $T_x = 709.2$. Then

$$x = \frac{\left\{ \begin{aligned} &(20)(245.7) - (8)(721.9) + (30)(56.5) \\ &+ (6)(2156.1) - (10)(546.2) - (10)(709.2) \end{aligned} \right\}}{32} = 38.0$$

The exact formulae for t -tests are complicated, and it appears that any good approximate rule would also be rather complicated. The following rule is suggested, though it somewhat underestimates the standard errors for the smaller squares and also lacks the simplicity that might be desired. For a comparison between the means of two treatments A and B , we assign to each an effective amount of replication which depends on the number and situation of the missing values. When scoring A we examine each replicate in turn and assign a score by the following rules.

Case	Score to A
I. A missing	0
II. A present, and B in the same row or column as A	
i. B missing	0
ii. B present	1
III. A present, and B not in the same row or column as A	
i. Other values missing in both row and column	$\frac{1}{2}$
ii. Other values missing in row or column but not in both	$\frac{2}{3}$
iii. No values missing in row or column	1

The effective replication for A is of course its total score over all replications in the experiment. B is scored similarly. Suppose that in the example the mean of treatment (18) were being compared with that of treatment (9), which occurs in the same row in square I. The effective replication for treatment (18) is 2, since case I arises in square I. The effective replication for treatment (9) is also 2, since case II(i) arises in square I.

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PLANS

Plan 12.1

3×3 balanced lattice square

$t = 9, k = 3, r = 2$, rows = 6, columns = 6, $\lambda = 1^*$

Square I

1	2	3
4	5	6
7	8	9

Square II

1	6	8
9	2	4
5	7	3

Plan 12.2

4×4 balanced lattice square

$t = 16, k = 4, r = 5$, rows = 20, columns = 20, $\lambda = 2$

Square I

1	5	9	13
2	6	10	14
3	7	11	15
4	8	12	16

Square II

1	2	3	4
6	5	8	7
11	12	9	10
16	15	14	13

Square III

1	11	16	6
12	2	5	15
14	8	3	9
7	13	10	4

Square IV

1	7	12	14
8	2	13	11
10	16	3	5
15	9	6	4

Square V

1	10	15	8
9	2	7	16
13	6	3	12
5	14	11	4

* Number of times that two treatments appear in the same row or column.

Plan 12.3

 5×5 balanced lattice square $t = 25, k = 5, r = 3, \text{rows} = 15, \text{columns} = 15, \lambda = 1$

Square I				
1	2	3	4	5
6	7	8	9	10
11	12	13	14	15
16	17	18	19	20
21	22	23	24	25

Square II				
1	10	14	18	22
23	2	6	15	19
20	24	3	7	11
12	16	25	4	8
9	13	17	21	5

Square III				
1	8	15	17	24
25	2	9	11	18
19	21	3	10	12
13	20	22	4	6
7	14	16	23	5

Plan 12.4

 7×7 balanced lattice square $t = 49, k = 7, r = 4, \text{rows} = 28, \text{columns} = 28, \lambda = 1$

Square I						
1	2	3	4	5	6	7
8	9	10	11	12	13	14
15	16	17	18	19	20	21
22	23	24	25	26	27	28
29	30	31	32	33	34	35
36	37	38	39	40	41	42
43	44	45	46	47	48	49

Square II						
1	38	26	14	44	32	20
21	2	39	27	8	45	33
34	15	3	40	28	9	46
47	35	16	4	41	22	10
11	48	29	17	5	42	23
24	12	49	30	18	6	36
37	25	13	43	31	19	7

Square III						
1	19	30	48	10	28	39
40	2	20	31	49	11	22
23	41	3	21	32	43	12
13	24	42	4	15	33	44
45	14	25	36	5	16	34
35	46	8	26	37	6	17
18	29	47	9	27	38	7

Square IV						
1	42	27	12	46	31	16
17	2	36	28	13	47	32
33	18	3	37	22	14	48
49	34	19	4	38	23	8
9	43	35	20	5	39	24
25	10	44	29	21	6	40
41	26	11	45	30	15	7

Plan 12.5

 8×8 balanced lattice square $t = 64, k = 8, r = 9, \text{rows} = 72, \text{columns} = 72, \lambda = 2$

Square I

1	9	17	25	33	41	49	57
2	10	18	26	34	42	50	58
3	11	19	27	35	43	51	59
4	12	20	28	36	44	52	60
5	13	21	29	37	45	53	61
6	14	22	30	38	46	54	62
7	15	23	31	39	47	55	63
8	16	24	32	40	48	56	64

Square II

1	10	19	28	37	46	55	64
9	2	51	44	61	30	23	40
17	50	3	36	29	62	15	48
25	42	35	4	21	14	63	56
33	58	27	20	5	54	47	16
41	26	59	12	53	6	39	24
49	18	11	60	45	38	7	32
57	34	43	52	13	22	31	8

Square III

1	44	62	56	27	39	18	13
46	2	17	35	16	53	60	31
64	23	3	25	54	12	45	34
55	40	29	4	58	41	11	22
28	9	50	63	5	24	38	43
37	51	15	42	20	6	32	57
19	61	48	14	33	26	7	52
10	30	36	21	47	59	49	8

Square IV

1	60	54	40	43	15	26	21
62	2	25	11	24	37	52	47
56	31	3	41	38	20	61	10
39	16	45	4	50	57	19	30
44	17	34	55	5	32	14	59
13	35	23	58	28	6	48	49
27	53	64	22	9	42	7	36
18	46	12	29	63	51	33	8

Square V

1	11	20	30	34	48	53	63
15	2	56	45	59	28	22	33
21	52	3	39	32	58	9	46
26	47	38	4	17	13	64	51
40	62	31	19	5	49	42	12
43	25	61	16	55	6	36	18
54	24	10	57	44	35	7	29
60	37	41	50	14	23	27	8

Square VI

1	59	52	38	42	16	29	23
63	2	32	13	19	36	54	41
53	28	3	47	40	18	57	14
34	15	46	4	49	61	24	27
48	22	39	51	5	25	10	60
11	33	21	64	31	6	44	50
30	56	58	17	12	43	7	37
20	45	9	26	62	55	35	8

Square VII

1	32	47	61	22	50	12	35
29	2	14	49	39	64	43	20
42	13	3	24	60	33	30	55
59	54	18	4	48	31	37	9
23	36	57	46	5	11	56	26
52	63	40	27	10	6	17	45
16	41	28	34	51	21	7	62
38	19	53	15	25	44	58	8

Square VIII

1	14	24	31	36	45	51	58
12	2	55	48	57	27	21	38
22	49	3	37	26	63	16	44
32	43	33	4	23	10	62	53
35	64	30	18	5	52	41	15
47	29	60	9	56	6	34	19
50	20	13	59	46	40	7	25
61	39	42	54	11	17	28	8

Square IX

1	2	3	4	5	6	7	8
14	12	16	10	15	9	13	11
24	21	22	23	18	19	20	17
31	27	26	32	30	29	25	28
36	38	37	33	35	34	40	39
45	48	44	43	41	47	46	42
51	55	49	53	52	56	50	54
58	57	63	62	64	60	59	61

Plan 12.8

 9×9 balanced lattice square $t = 81, k = 9, r = 5, \text{rows} = 45, \text{columns} = 45, \lambda = 1$

Square I

1	2	3	4	5	6	7	8	9
10	11	12	13	14	15	16	17	18
19	20	21	22	23	24	25	26	27
28	29	30	31	32	33	34	35	36
37	38	39	40	41	42	43	44	45
46	47	48	49	50	51	52	53	54
55	56	57	58	59	60	61	62	63
64	65	66	67	68	69	70	71	72
73	74	75	76	77	78	79	80	81

Square II

1	12	20	34	45	53	58	69	77
21	2	10	54	35	43	78	59	67
11	19	3	44	52	36	68	76	60
61	72	80	4	15	23	28	39	47
81	62	70	24	5	13	48	29	37
71	79	63	14	22	6	38	46	30
31	42	50	55	66	74	7	18	26
51	32	40	75	56	64	27	8	16
41	49	33	65	73	57	17	25	9

Square III

1	57	29	16	72	44	22	78	50
30	2	55	45	17	70	51	23	76
56	28	3	71	43	18	77	49	24
25	81	53	4	60	32	10	66	38
54	26	79	33	5	58	39	11	64
80	52	27	59	31	6	65	37	12
13	69	41	19	75	47	7	63	35
42	14	67	48	20	73	36	8	61
68	40	15	74	46	21	62	34	9

Square IV

1	33	62	27	47	76	14	43	66
63	2	31	77	25	48	64	15	44
32	61	3	46	78	26	45	65	13
17	37	69	4	36	56	21	50	79
67	18	38	57	5	34	80	19	51
39	68	16	35	55	6	49	81	20
24	53	73	11	40	72	7	30	59
74	22	54	70	12	41	60	8	28
52	75	23	42	71	10	29	58	9

Square V

1	60	35	18	65	40	23	79	48
36	2	58	41	16	66	46	24	80
59	34	3	64	42	17	81	47	22
26	73	51	4	63	29	12	68	43
49	27	74	30	5	61	44	10	69
75	50	25	62	28	6	67	45	11
15	71	37	20	76	54	7	57	32
38	13	72	52	21	77	33	8	55
70	39	14	78	53	19	56	31	9

LATTICE SQUARES

Plan 12.7

 11×11 balanced lattice square $t = 121, k = 11, r = 6, \text{rows} = 66, \text{columns} = 66, \lambda = 1$

Square I

1	2	3	4	5	6	7	8	9	10	11
12	13	14	15	16	17	18	19	20	21	22
23	24	25	26	27	28	29	30	31	32	33
34	35	36	37	38	39	40	41	42	43	44
45	46	47	48	49	50	51	52	53	54	55
56	57	58	59	60	61	62	63	64	65	66
67	68	69	70	71	72	73	74	75	76	77
78	79	80	81	82	83	84	85	86	87	88
89	90	91	92	93	94	95	96	97	98	99
100	101	102	103	104	105	106	107	108	109	110
111	112	113	114	115	116	117	118	119	120	121

Square II

1	102	82	62	42	22	112	92	72	52	32
33	2	103	83	63	43	113	93	73	53	
54	23	3	104	84	64	114	94	74		
75	55	24	4	105	85	65	34	14	115	95
96	76	45	25	5	106	86	66	35	15	116
117	97	77	46	26	6	107	87	56	36	16
17	118	98	67	47	27	7	108	88	57	37
38	18	119	99	68	48	28	8	109	78	58
59	39	19	120	89	69	49	29	9	110	79
80	60	40	20	121	90	70	50	30	10	100
101	81	61	41	21	111	91	71	51	31	11

Square III

1	115	108	90	83	76	58	51	44	26	19
20	2	116	109	91	84	77	59	52	34	27
28	21	3	117	110	92	85	67	60	53	35
36	29	22	4	118	100	93	86	68	61	54
55	37	30	12	5	119	101	94	87	69	62
63	45	38	31	13	6	120	102	95	88	70
71	64	46	39	32	14	7	121	103	96	78
79	72	65	47	40	33	15	8	111	104	97
98	80	73	66	48	41	23	16	9	112	105
106	99	81	74	56	49	42	24	17	10	113
114	107	89	82	75	57	50	43	25	18	11

Plan 12.7 (Continued) 11×11 balanced lattice square

Square IV

1	40	68	107	14	53	81	120	27	66	94
95	2	41	69	108	15	54	82	121	28	56
57	96	3	42	70	109	16	55	83	111	29
30	58	97	4	43	71	110	17	45	84	112
113	31	59	98	5	44	72	100	18	46	85
86	114	32	60	99	6	34	73	101	19	47
48	87	115	33	61	89	7	35	74	102	20
21	49	88	116	23	62	90	8	36	75	103
104	22	50	78	117	24	63	91	9	37	76
77	105	12	51	79	118	25	64	92	10	38
39	67	106	13	52	80	119	26	65	93	11

Square V

1	119	105	91	88	74	60	46	43	29	15
16	2	120	106	92	78	75	61	47	44	30
31	17	3	121	107	93	79	76	62	48	34
35	32	18	4	111	108	94	80	77	63	49
50	36	33	19	5	112	109	95	81	67	64
65	51	37	23	20	6	113	110	96	82	68
69	66	52	38	24	21	7	114	100	97	83
84	70	56	53	39	25	22	8	115	101	98
99	85	71	57	54	40	26	12	9	116	102
103	89	86	72	58	55	41	27	13	10	117
118	104	90	87	73	59	45	42	28	14	11

Square VI

1	110	87	64	41	18	116	93	70	47	24
25	2	100	88	65	42	19	117	94	71	48
49	26	3	101	78	66	43	20	118	95	72
73	50	27	4	102	79	56	44	21	119	96
97	74	51	28	5	103	80	57	34	22	120
121	98	75	52	29	6	104	81	58	35	12
13	111	99	76	53	30	7	105	82	59	36
37	14	112	89	77	54	31	8	106	83	60
61	38	15	113	90	67	55	32	9	107	84
85	62	39	16	114	91	68	45	33	10	108
109	86	63	40	17	115	92	69	46	23	11

Plan 12.8

 13×13 balanced lattice square $t = 169, k = 13, r = 7, \text{rows} = 91, \text{columns} = 91, \lambda = 1$

Square I

1	2	3	4	5	6	7	8	9	10	11	12	13
14	15	16	17	18	19	20	21	22	23	24	25	26
27	28	29	30	31	32	33	34	35	36	37	38	39
40	41	42	43	44	45	46	47	48	49	50	51	52
53	54	55	56	57	58	59	60	61	62	63	64	65
66	67	68	69	70	71	72	73	74	75	76	77	78
79	80	81	82	83	84	85	86	87	88	89	90	91
92	93	94	95	96	97	98	99	100	101	102	103	104
105	106	107	108	109	110	111	112	113	114	115	116	117
118	119	120	121	122	123	124	125	126	127	128	129	130
131	132	133	134	135	136	137	138	139	140	141	142	143
144	145	146	147	148	149	150	151	152	153	154	155	156
157	158	159	160	161	162	163	164	165	166	167	168	169

Square II

1	26	38	50	62	74	86	98	110	122	134	146	158
159	2	14	39	51	63	75	87	99	111	123	135	147
148	160	3	15	27	52	64	76	88	100	112	124	136
137	149	161	4	16	28	40	65	77	89	101	113	125
126	138	150	162	5	17	29	41	53	78	90	102	114
115	127	139	151	163	6	18	30	42	54	66	91	103
104	116	128	140	152	164	7	19	31	43	55	67	79
80	92	117	129	141	153	165	8	20	32	44	56	68
69	81	93	105	130	142	154	166	9	21	33	45	57
58	70	82	94	106	118	143	155	167	10	22	34	46
47	59	71	83	95	107	119	131	156	168	11	23	35
36	48	60	72	84	96	108	120	132	144	169	12	24
25	37	49	61	73	85	97	109	121	133	145	157	13

Plan 12.8 (Continued) 13×13 balanced lattice square

Square III

1	24	34	44	54	77	87	97	107	130	140	150	160
161	2	25	35	45	55	78	88	98	108	118	141	151
152	162	3	26	36	46	56	66	89	99	109	119	142
143	153	163	4	14	37	47	57	67	90	100	110	120
121	131	154	164	5	15	38	48	58	68	91	101	111
112	122	132	155	165	6	16	39	49	59	69	79	102
103	113	123	133	156	166	7	17	27	50	60	70	80
81	104	114	124	134	144	167	8	18	28	51	61	71
72	82	92	115	125	135	145	168	9	19	29	52	62
63	73	83	93	116	126	136	146	169	10	20	30	40
41	64	74	84	94	117	127	137	147	157	11	21	31
32	42	65	75	85	95	105	128	138	148	158	12	22
23	33	43	53	76	86	96	106	129	139	149	159	13

Square IV

1	22	30	51	59	67	88	96	117	125	133	154	162
163	2	23	31	52	60	68	89	97	105	126	134	155
156	164	3	24	32	40	61	69	90	98	106	127	135
136	144	165	4	25	33	41	62	70	91	99	107	128
129	137	145	166	5	26	34	42	63	71	79	100	108
109	130	138	146	167	6	14	35	43	64	72	80	101
102	110	118	139	147	168	7	15	36	44	65	73	81
82	103	111	119	140	148	169	8	16	37	45	53	74
75	83	104	112	120	141	149	157	9	17	38	46	54
55	76	84	92	113	121	142	150	158	10	18	39	47
48	56	77	85	93	114	122	143	151	159	11	19	27
28	49	57	78	86	94	115	123	131	152	160	12	20
21	29	50	58	66	87	95	116	124	132	153	161	13

Plan 12.8 (Continued) 13×13 balanced lattice square

Square V												
1	20	39	45	64	70	89	95	114	120	139	145	164
165	2	21	27	46	65	71	90	96	115	121	140	146
147	166	3	22	28	47	53	72	91	97	116	122	141
142	148	167	4	23	29	48	54	73	79	98	117	123
124	143	149	168	5	24	30	49	55	74	80	99	105
106	125	131	150	169	6	25	31	50	56	75	81	100
101	107	126	132	151	157	7	26	32	51	57	76	82
83	102	108	127	133	152	158	8	14	33	52	58	77
78	84	103	109	128	134	153	159	9	15	34	40	59
60	66	85	104	110	129	135	154	160	10	16	35	41
42	61	67	86	92	111	130	136	155	161	11	17	36
37	43	62	68	87	93	112	118	137	156	162	12	18
19	38	44	63	69	88	94	113	119	138	144	163	13

Square VI												
1	18	35	52	56	73	90	94	111	128	132	149	166
167	2	19	36	40	57	74	91	95	112	129	133	150
151	168	3	20	37	41	58	75	79	96	113	130	134
135	152	169	4	21	38	42	59	76	80	97	114	118
119	136	153	157	5	22	39	43	60	77	81	98	115
116	120	137	154	158	6	23	27	44	61	78	82	99
100	117	121	138	155	159	7	24	28	45	62	66	83
84	101	105	122	139	156	160	8	25	29	46	63	67
68	85	102	106	123	140	144	161	9	26	30	47	64
65	69	86	103	107	124	141	145	162	10	14	31	48
49	53	70	87	104	108	125	142	146	163	11	15	32
33	50	54	71	88	92	109	126	143	147	164	12	16
17	34	51	55	72	89	93	110	127	131	148	165	13

Plan 12.8 (Continued) 13×13 balanced lattice square

Square VII

1	16	31	46	61	76	91	93	108	123	138	153	168
169	2	17	32	47	62	77	79	94	109	124	139	154
155	157	3	18	33	48	63	78	80	95	110	125	140
141	156	158	4	19	34	49	64	66	81	96	111	126
127	142	144	159	5	20	35	50	65	67	82	97	112
113	128	143	145	160	6	21	36	51	53	68	83	98
99	114	129	131	146	161	7	22	37	52	54	69	84
85	100	115	130	132	147	162	8	23	38	40	55	70
71	86	101	116	118	133	148	163	9	24	39	41	56
57	72	87	102	117	119	134	149	164	10	25	27	42
43	58	73	88	103	105	120	135	150	165	11	26	28
29	44	59	74	89	104	106	121	136	151	166	12	14
15	30	45	60	75	90	92	107	122	137	152	167	13

CHAPTER 13

INCOMPLETE LATIN SQUARES (YOUTDEN SQUARES)

13.1 Description

13.11 Youden Squares. These designs, which are constructed by a rearrangement of certain of the balanced incomplete blocks, possess the characteristic "double control" of the latin square, without the restriction that the number of replicates must equal the number of treatments. A latin square with 13 treatments is rarely used, because it necessitates 13 replications. There are, however, incomplete latin squares for 13 treatments in either 4 or 9 replicates. Every treatment occurs once in a column (replication), and every pair of treatments appears together an equal number of times in the same block. Column differences are eliminated automatically from the treatment comparisons, while block differences may be removed by adjusting the treatment mean yields in the same way as with balanced incomplete blocks. Most of the designs were developed by Youden (13.1, 13.2), whose name is commonly associated with them; previously Yates (13.3) had drawn attention to the group of designs in which the number of replicates is one less than the number of treatments. Table 13.1 summarizes the designs for numbers of treatments up to 37.

TABLE 13.1 SUMMARY OF INCOMPLETE LATIN SQUARES (r less than t)

t	$r = k$	λ	E	Type †	t	$r = k$	λ	E	Type †
4 *	3	2	89	II	13	4	1	81	I
5 *	4	3	94	II	13	9	6	95	I
6 *	5	4	96	II	15	7	3	92	I
7	3	1	78	II	15	8	4	94	I
7	4	2	88	II	16	6	2	89	I
7 *	6	5	97	II	16	10	6	96	I
8 *	7	6	98	II	19	9	4	94	I
9 *	8	7	98	II	19	10	5	95	I
10 *	9	8	99	II	21	5	1	84	I
11	5	2	88	I	31	6	1	86	I
11	6	3	92	I	37	9	2	91	I
11 *	10	9	99	II					

* Constructed from a $t \times t$ latin square by omission of the last column.

† This refers to the method of analysis (see section 13.2).

The number of replicates is always equal to the number of units per block. The symbol λ shows the number of times that two treatments appear together in a block, while E is the efficiency factor expressed in percentage.

Plans are given (plans 13.1–13.15) for all designs in table 13.1 except those marked with an asterisk (*), which are constructed from latin squares by the omission of the last column. A 5×5 square, for instance, furnishes the plan for $t = 5, r = 4$.

An interesting application of the design for the control of plant variability was made by Youden (13.1) in greenhouse experiments on tobacco-mosaic virus. The experimental unit was a single leaf, and the data consisted of the number of lesions produced per leaf by rubbing the leaf with a solution which contained the virus. The numbers of lesions had been found to depend much more on inherent qualities of the plant than on the position of the plant on the greenhouse bench. Consequently, each block of the design was a single plant, so that the large differences in responsiveness which existed among plants did not contribute to the experimental errors. The columns were the positions, from top to bottom, of the five leaves which were used on each plant. That is, the first replication contained the top leaf of every plant. Since there was a fairly consistent gradient in responsiveness down each plant, this control also proved effective.

As in the example above, when laying out an incomplete latin square the general principle is to group the units so that differences among blocks and differences among columns represent the major sources of variation that are known or suspected.

13.12 Randomization. The steps are

1. Rearrange the blocks of the plan at random.
2. Rearrange the replications of the plan at random.

When a plan is repeated for additional replication, blocks and columns are randomized separately within each *repetition*. The repetitions may be kept separate.

13.2 Statistical Analysis

In experiments with more than 10 treatments, the numbers of degrees of freedom for blocks are large enough to allow the use of inter-block information. The appropriate analysis is described in the next section. For small experiments, where inter-block information should be ignored, the analysis is given in section 13.22. The recommended method of analysis for any experiment is indicated both in table 13.1 and in the plans by the "type" symbol (I or II).

13.21 Type I. Analysis with Recovery of Inter-block Information. The steps are as follows (t = number of treatments, k = number of units per block = r).

1. Calculate the treatment totals T , the column totals, the block totals, and the grand total G .

2. For each treatment, calculate the total B_t of all the blocks which contain the treatment. Place these quantities in a column next to the column of treatment totals. Then form a third column of the quantities:

$$W = (t - k)T - (t - 1)B_t + (k - 1)G$$

The W 's should sum to zero.

3. The separation of degrees of freedom in the analysis of variance is as follows.

	d.f.	m.s.
Columns (replications)	$(k - 1)$	
Blocks (eliminating treatments)	$(t - 1)$	E_b
Treatments	$(t - 1)$	
Error	$(k - 2)(t - 1)$	E_e
Total	$(tk - 1)$	

The total s.s. and the sums of squares for columns and treatments are found by the usual methods. The sum of squares for blocks (eliminating treatments) is the sum of squares of the W 's, divided by $kt(t - k)(k - 1)$. The error s.s. is obtained by subtraction.

4. Calculate the factor

$$\mu = \frac{(E_b - E_e)}{t(k - 1)E_b}$$

The adjusted total for any treatment is

$$Y = T + \mu W$$

Should E_b be less than E_e , μ is taken as zero and no adjustments are made to the treatment totals. The adjusted treatment means are obtained on dividing each Y by r . The estimated error variance of the difference between two adjusted treatment means is

$$\frac{2E_e}{r} [1 + (t - r)\mu]$$

13.22 Type II. Analysis without Recovery of Inter-block Information. The analysis proceeds as follows.

1. Calculate the column totals, the block totals, the treatment totals T , and the grand total G .

2. For each treatment, obtain the quantity

$$Q = kT - B_i$$

where B_i is the total of all the blocks in which the treatment appears. The quantities Q should sum to zero.

3. The degrees of freedom in the analysis of variance are partitioned as in section 13.21. In this case the blocks s.s. is calculated without eliminating treatment effects, while the treatments s.s. is adjusted for block differences. The total s.s. and the sums of squares for blocks are computed by the standard procedure. The treatments s.s. (adjusted) is the sum of the squares of the Q 's, divided by $tk\lambda$. The error s.s. is found by subtraction.

4. In order to obtain any treatment mean, adjusted for block differences, divide the corresponding Q by $t\lambda$ and add to the quotient the mean for the whole experiment.

The estimated error variance of the difference between two adjusted treatment means is

$$\frac{2E_e(t-1)}{(r-1)t}$$

Numerical examples of this analysis for $t = 7$, $r = 3$, and $t = 21$, $r = 5$ are given in reference (13.1), and an example for $t = 6$, $r = 5$ in (13.3).

13.23 Type Ia. Repetitions of Type I. Let n be the number of replications in the basic plan, which is used p times in an experiment, so that the total number of replications $r = np$.

The method of analysis presented here may be used with *any* of the designs in table 13.1, since even with small numbers of treatments the repetition provides sufficient degrees of freedom for estimating the inter-block variation. If, however, the efficiency factor exceeds 95%, it is scarcely worth while to utilize inter-block information.

The changes necessary in the procedure of section 13.21 are outlined below, the numbers referring to the steps in that section.

1. Unchanged. Corresponding to any block, there are $(p-1)$ other blocks which contain the same set of treatments. The block totals should be arranged in a table (table A , say) with t rows and p columns.

2. Unchanged. Note that the calculation of the quantities B_i is expedited by the use of the row totals of table A .

3. The analysis of variance is as follows.

	d.f.	m.s.
Columns (replications)	$(kp - 1)$	
Blocks		
Component (a)	$(p - 1)(t - 1)$	
Component (b)	$(t - 1)$	
Block total	$p(t - 1)$	E_b
Treatments	$(t - 1)$	
Error	$(t - 1)(pk - p - 1)$	E_e
Total	$(pkt - 1)$	

In this analysis, the $(p - 1)$ degrees of freedom among repetitions have been ascribed to the columns, so that the blocks s.s. is actually a sum of squares among blocks *within repetitions*.

Component (a) of the sum of squares for blocks consists of comparisons among blocks which contain the same set of treatments and is the interaction s.s. for table A.

Component (b) is the sum of squares of the W 's, with a divisor $pkt(t - k)(k - 1)$.

4. Unchanged except that

$$\mu = \frac{p(E_b - E_e)}{pt(k - 1)E_b - (t - k)(p - 1)E_e}$$

The estimated error variance of the difference between two adjusted treatment means is

$$\frac{2E_e}{r} [1 + (t - n)\mu]$$

13.24 Type IIa. Repetitions of Type II. As we have pointed out, in designs where the efficiency factor is high, inter-block information can be neglected. The changes required from section 13.22 should present no difficulty. The degrees of freedom subdivide as in section 13.23. The $p(t - 1)$ degrees of freedom among blocks within repetitions are calculated in the usual way, each repetition furnishing $(t - 1)$ degrees of freedom. Notice that the divisors for the treatments s.s. and for the quantities Q must be increased by the factor p .

13.25 Missing Data. The formula for inserting an estimate in place of a missing observation is obtained by minimizing the intra-block error. This is the correct estimate for the smaller experiments where inter-block information is ignored. For reasons given in section 10.13, this

estimate will also be used to provide an approximate solution when inter-block information is recovered. The formula is quite similar to that for balanced incomplete blocks, with the addition of an extra term involving the column (or replication) total.

Let C , B , and T be the totals of the column, block, and treatment that contain the missing value, and let G be the grand total. Further, as in the statistical analysis, let B_t be the total of all blocks in which the treatment with the missing value appears (there will, of course, be r such blocks). Finally,

T' = total (over all replicates) of all *other* treatments that appear in the block which has the missing value

B_t' = total of the B_t values for all *other* treatments that appear in the block with the missing value

The estimate x of the missing value is

$$x = \frac{\lambda[rC + tB + (t-1)T - G] - rT' - (r-1)B_t + B_t'}{r(r-1)(r-2)}$$

The symbol λ is given for each design in table 13.1. Care must be taken not to confuse the block totals with the column totals.

If the experiment contains p repetitions, the formula is

$$x = \frac{p[pk\lambda C + pt\lambda B + k(k-1)T - p\lambda R - kT' - (k-1)B_t + B_t']}{r(k-1)(pk-p-1)}$$

where R is the total of the repetition in which the missing value occurs, and λ is as in table 13.1.

13.3 Other Designs for Small Numbers of Treatments

13.31 Description. When the number of treatments is small, it is sometimes useful to have a design of the "latin square" type in which the number of replicates exceeds the number of treatments. Some designs can be obtained by repetition of an ordinary latin square or of a Youden square. Additional plans can be constructed by adding a Youden square to a latin square. A selection of these designs is shown in table 13.2 for numbers of treatments up to 7.

It will be noted that plans are available for most numbers of replicates up to 10. Actually, plans can be made for any number of replicates up to 10; those shown here have been selected for ease of analysis.

TABLE 13.2 INCOMPLETE LATIN SQUARES FOR SMALL NUMBERS OF TREATMENTS
($r > t$)

t	k	r	E	Ref. to plan	Type †	t	k	r	E	Ref. to plan	Type †
3	5	5	96	13.16	III	4	9	9	99	13.22	IV
3	3	6	100	2 L.S.		4	5	10	96	13.20	IIa
3	7	7	98	13.17	IV	5	6	6	97	13.23	IV
3	8	8	98	13.18	III	5	4	8	94	*	Ia
3	3	9	100	3 L.S.		5	9	9	99	13.24	III
3	5	10	96	13.16	IIa	5	5	10	100	2 L.S.	
3	10	10	99	13.19	IV	6	7	7	98	13.25	IV
4	5	5	96	13.20	IV	6	5	10	96	*	Ia
4	3	6	89	*	Ia	7	4	8	88	13.2	Ia
4	7	7	98	13.21	III	7	8	8	98	13.26	IV
4	4	8	100	2 L.S.		7	3	9	78	13.1	Ia
4	3	9	89	*	Ia						

* By repetition of the plan for $r = t - 1$, which is constructed by taking a $t \times t$ latin square and omitting the last column.

† This refers to the method of analysis (see section 13.2).

TABLE 13.3 INCOMPLETE LATIN SQUARE FOR 4 TREATMENTS IN 7 REPLICATES

Treatment symbols and yields of tomatoes (pounds)

Block	Column							Total
	I	II	III	IV	V	VI	VII	
1	2	2	4	4	3	3	1	
	50	72	83	82	76	89	74	526
2	1	3	3	1	4	2	4	
	40	59	71	91	59	73	52	445
3	4	1	2	3	1	4	2	
	43	57	58	98	54	71	51	432
4	3	4	1	2	2	1	3	
	48	54	74	97	75	75	54	477
Totals	181	242	286	368	264	308	231	1880

	T	$Q = kT + B'$	$Q \div 48$	Adjusted means (subtract 11.2)
1	465	3,781	78.8	67.6
2	476	3,777	78.7	67.5
3	495	3,897	81.2	70.0
4	444	3,585	74.7	63.5
Check totals or mean	1,880	15,040		67.1

13.32 Type III. Statistical Analysis When $k = it - 1$. In this case the size of block, which equals the number of replicates, is 1 less than some multiple (i) of the number of treatments. Since these designs have not been discussed in the literature, an example is given from uniformity data. Table 13.3 shows the tomato yields in pounds of 28 single-row plots, Hartman and Stair's data (13.4), on which the design for 4 treatments in 7 replicates has been superimposed.

Since the plots measured 6 feet \times 24 feet, each column (replication) is compact, being 24 feet square. There are obviously substantial differences among replicates. The blocks, although very oblong, may also exhibit differences in yield, because each block is a separate row of plants.

The efficiency factors of these designs are all very high, so that inter-block information is ignored in the analysis.

The steps in the analysis are as follows:

1. Calculate the column totals, the block totals, the treatment totals T , and the grand total G .

2. A property of these designs is that each treatment is replicated less in one block than in the other blocks. Thus, in the example, treatment (1) appears only once in the first block, but twice in all other blocks. Similarly, treatment (2) is deficient in block 2 and so on. For each treatment calculate the quantity

$$Q = kT + B'$$

where B' is the total for the block in which the treatment is deficient. For instance

$$Q_1 = 7 \times 465 + 526 = 3781$$

As a check, the quantities Q should sum to $(k + 1)G$.

3. All sums of squares in the analysis of variance are obtained in the usual way except that for treatments, which is given by the sum of squares of deviations of the Q 's, divided by $k(k^2 - 1)$. In this case we have

$$\frac{(3781)^2 + (3777)^2 + (3897)^2 + (3585)^2 - \frac{1}{4}(15,040)^2}{336} = 150$$

The analysis of variance is shown below. The degrees of freedom subdivide as in section 13.21.

	d.f.	s.s.	m.s.
Columns (replications)	6	5387	897.8
Blocks	3	750	250.0
Treatments	3	150	50.0
Error	15	766	51.1
Total	27	7053	

The elimination of block differences has substantially reduced the error m.s.

4. To obtain the adjusted treatment mean yields, we first divide each Q by $(k^2 - 1)$, in this case 48. From the resulting quantities we subtract $G/tk(k - 1)$, or $1880/168 = 11.2$. As a further check, the mean of the adjusted treatment means should equal the mean yield of the whole experiment (67.1). The efficiency factor is $(k^2 - 1)/k^2$, and the estimated error variance of the difference between two adjusted treatment means is

$$\frac{2E_e r}{r^2 - 1}$$

13.33 Type IV. Statistical Analysis When $k = it + 1$. The analysis closely resembles that of section 13.32. For each treatment there is one block in which the treatment has extra replication. Thus, in plan 13.17 for $t = 3$, $r = 7$, treatment (1) appears 3 times in the first block but only twice in any other block. The only changes in the computing instructions of section 13.32 are (i) $Q = kT - B'$, where B' is the total of the block in which the treatment has extra replication, (ii) the Q 's sum to $(k - 1)G$, and (iii) for the adjusted treatment means, divide Q by $(k^2 - 1)$, and add to the quotient the quantity $G/tk(k - 1)$. The change in the divisor of G should be noted.

The error variance of the difference between two adjusted means remains as in section 13.24.

REFERENCES

- 13.1 YODEN, W. J. Use of incomplete block replications in estimating tobacco-mosaic virus. *Contr. Boyce Thompson Inst.* 9, 41-48, 1937.
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- 13.3 YATES, F. Incomplete latin squares. *Jour. Agr. Sci.* 26, 301-315, 1936.
- 13.4 HARTMAN, J. D., and STAIR, E. C. Field plot technique studies with tomatoes. *Proc. Amer. Soc. Hort. Sci.* 41, 315-320, 1942.

PLANS

Plan 13.1 $t = 7, k = 3, r = 3, b = 7, \lambda = 1, E = 78$, Type II

Reps.			
Block	I	II	III
(1)	7	1	3
(2)	1	2	4
(3)	2	3	5
(4)	3	4	6
(5)	4	5	7
(6)	5	6	1
(7)	6	7	2

Plan 13.2 $t = 7, k = 4, r = 4, b = 7, \lambda = 2, E = 88$, Type II

Reps.				
Block	I	II	III	IV
(1)	3	5	6	7
(2)	4	6	7	1
(3)	5	7	1	2
(4)	6	1	2	3
(5)	7	2	3	4
(6)	1	3	4	5
(7)	2	4	5	6

Plan 13.3 $t = 11, k = 5, r = 5, b = 11, \lambda = 2, E = 88$, Type I

Reps.					
Block	I	II	III	IV	V
(1)	1	2	3	4	5
(2)	7	1	6	10	3
(3)	9	8	1	6	2
(4)	11	9	7	1	4
(5)	10	11	5	8	1
(6)	8	7	2	3	11
(7)	2	6	4	11	10
(8)	6	3	11	5	9
(9)	3	4	10	9	8
(10)	5	10	9	2	7
(11)	4	5	8	7	6

Plan 13.4 $t = 11, k = 6, r = 6, b = 11, \lambda = 3, E = 92$, Type I

Block	Reps.					
	I	II	III	IV	V	VI
(1)	6	7	8	9	10	11
(2)	5	8	4	11	2	9
(3)	4	5	7	3	11	10
(4)	3	10	2	6	5	8
(5)	2	3	9	7	4	6
(6)	1	6	10	4	9	5
(7)	9	1	3	5	8	7
(8)	8	2	1	10	7	4
(9)	7	11	5	1	6	2
(10)	11	4	6	8	1	3
(11)	10	9	11	2	3	1

Plan 13.5 $t = 13, k = 4, r = 4, b = 13, \lambda = 1, E = 81$, Type I

Block	Reps.			
	I	II	III	IV
(1)	13	1	3	9
(2)	1	2	4	10
(3)	2	3	5	11
(4)	3	4	6	12
(5)	4	5	7	13
(6)	5	6	8	1
(7)	6	7	9	2
(8)	7	8	10	3
(9)	8	9	11	4
(10)	9	10	12	5
(11)	10	11	13	6
(12)	11	12	1	7
(13)	12	13	2	8

Plan 13.6 $t = 13, k = 9, r = 9, b = 13, \lambda = 6, E = 95$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII	VIII	IX
(1)	2	5	6	7	9	10	11	12	13
(2)	3	6	7	8	10	11	12	13	1
(3)	4	7	8	9	11	12	13	1	2
(4)	5	8	9	10	12	13	1	2	3
(5)	6	9	10	11	13	1	2	3	4
(6)	7	10	11	12	1	2	3	4	5
(7)	8	11	12	13	2	3	4	5	6
(8)	9	12	13	1	3	4	5	6	7
(9)	10	13	1	2	4	5	6	7	8
(10)	11	1	2	3	5	6	7	8	9
(11)	12	2	3	4	6	7	8	9	10
(12)	13	3	4	5	7	8	9	10	11
(13)	1	4	5	6	8	9	10	11	12

Plan 13.7 $t = 15, k = 7, r = 7, b = 15, \lambda = 3, E = 92$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII
(1)	13	8	12	6	7	1	9
(2)	5	14	10	7	12	2	8
(3)	15	12	11	5	8	3	6
(4)	12	11	6	9	2	4	14
(5)	4	5	8	1	14	9	15
(6)	11	9	7	2	13	15	5
(7)	1	2	3	4	5	6	7
(8)	2	3	1	13	15	14	12
(9)	8	6	4	15	10	13	2
(10)	10	4	5	11	1	12	13
(11)	9	13	14	10	6	5	3
(12)	14	7	13	3	4	8	11
(13)	7	15	9	12	3	10	4
(14)	3	1	2	8	9	11	10
(15)	6	10	15	14	11	7	1

Plan 13.8 $t = 15, k = 8, r = 8, b = 15, \lambda = 4, E = 94$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII	VIII
(1)	11	4	2	5	10	3	14	15
(2)	4	1	3	15	13	11	6	9
(3)	9	2	14	4	7	1	10	13
(4)	15	3	1	10	8	13	7	5
(5)	7	13	10	12	11	2	3	6
(6)	6	10	12	1	4	14	8	3
(7)	12	9	13	14	15	10	11	8
(8)	10	11	9	7	6	8	5	4
(9)	5	14	7	3	1	9	12	11
(10)	8	15	6	2	3	7	9	14
(11)	1	7	11	8	2	4	15	12
(12)	2	6	15	9	5	12	1	10
(13)	13	8	5	11	14	6	2	1
(14)	14	5	4	6	12	15	13	7
(15)	3	12	8	13	9	5	4	2

Plan 13.9 $t = 16, k = 6, r = 6, b = 16, \lambda = 2, E = 89$, Type I

Reps.

Block	I	II	III	IV	V	VI
(1)	1	2	3	4	5	6
(2)	2	7	8	9	10	1
(3)	3	1	13	7	11	12
(4)	4	8	1	11	14	15
(5)	5	12	14	1	16	9
(6)	6	10	15	13	1	16
(7)	7	14	2	16	15	3
(8)	8	16	12	2	4	13
(9)	9	15	11	5	13	2
(10)	10	11	6	12	2	14
(11)	11	4	16	3	9	10
(12)	12	3	10	15	8	5
(13)	13	6	9	14	3	8
(14)	14	13	5	10	7	4
(15)	15	9	4	6	12	7
(16)	16	5	7	8	6	11

Plan 13.10 $t = 16, k = 10, r = 10, b = 16, \lambda = 6, E = 96$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII	VIII	IX	X
(1)	8	7	9	10	11	12	13	14	15	16
(2)	3	4	5	13	16	11	12	6	14	15
(3)	9	2	4	5	6	8	10	15	16	14
(4)	2	6	3	7	5	10	9	16	12	13
(5)	6	8	2	3	4	7	15	10	13	11
(6)	4	5	14	8	3	9	7	2	11	12
(7)	1	10	11	4	13	5	6	12	8	9
(8)	5	1	15	6	14	3	11	9	7	10
(9)	16	3	1	14	12	6	4	8	10	7
(10)	7	13	16	1	15	4	3	5	9	8
(11)	12	14	7	15	1	2	8	13	6	5
(12)	14	11	13	16	9	1	2	7	4	6
(13)	10	15	12	2	7	16	1	11	5	4
(14)	11	12	6	9	8	15	16	1	2	3
(15)	13	16	8	11	10	14	5	3	1	2
(16)	15	9	10	12	2	13	14	4	3	1

Plan 13.11 $t = 19, k = 9, r = 9, b = 19, \lambda = 4, E = 94$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII	VIII	IX
(1)	1	2	3	4	5	6	7	8	9
(2)	14	1	2	3	4	12	11	10	13
(3)	17	16	1	2	15	10	6	5	11
(4)	12	13	16	1	2	7	18	15	8
(5)	9	10	12	16	1	3	17	19	7
(6)	8	11	13	19	17	1	3	6	18
(7)	7	18	11	14	16	19	1	4	5
(8)	6	9	17	12	14	18	15	1	4
(9)	5	8	9	10	13	15	19	14	1
(10)	13	5	14	9	18	2	16	3	17
(11)	10	6	7	15	19	14	2	18	3
(12)	18	17	5	8	12	4	10	2	19
(13)	19	4	15	7	9	17	13	11	2
(14)	2	19	6	11	8	9	14	16	12
(15)	3	15	10	18	11	8	4	9	16
(16)	4	3	19	5	6	16	12	13	15
(17)	15	7	8	17	3	11	5	12	14
(18)	16	14	4	6	7	13	8	17	10
(19)	11	12	18	13	10	5	9	7	6

Plan 13.12 $t = 19, k = 10, r = 10, b = 19, \lambda = 5, E = 95$, Type I

Block	Reps.									
	I	II	III	IV	V	VI	VII	VIII	IX	X
(1)	10	11	12	13	14	15	16	17	18	19
(2)	19	5	6	7	8	9	15	16	17	18
(3)	18	19	3	4	7	8	9	12	13	14
(4)	14	17	19	11	4	5	6	9	10	3
(5)	13	2	15	14	18	4	5	6	11	8
(6)	12	14	2	10	15	7	4	5	9	16
(7)	15	13	17	2	12	6	8	10	3	9
(8)	16	10	11	19	2	3	7	13	8	5
(9)	11	16	18	17	6	2	3	4	12	7
(10)	1	8	10	15	11	19	12	7	4	6
(11)	17	1	8	9	16	12	13	11	5	4
(12)	6	15	1	16	9	14	11	3	7	13
(13)	5	6	14	1	3	10	18	8	16	12
(14)	4	7	5	3	1	13	17	18	15	10
(15)	7	12	13	5	19	1	2	14	6	17
(16)	8	9	7	18	10	17	1	2	14	11
(17)	9	4	16	6	13	18	10	1	19	2
(18)	3	18	9	12	5	11	19	15	2	1
(19)	2	3	4	8	17	16	14	19	1	15

Plan 13.13 $t = 21, k = 5, r = 5, b = 21, \lambda = 1, E = 84$, Type I

Reps.

Block	I	II	III	IV	V
(1)	21	1	4	14	16
(2)	1	2	5	15	17
(3)	2	3	6	16	18
(4)	3	4	7	17	19
(5)	4	5	8	18	20
(6)	5	6	9	19	21
(7)	6	7	10	20	1
(8)	7	8	11	21	2
(9)	8	9	12	1	3
(10)	9	10	13	2	4
(11)	10	11	14	3	5
(12)	11	12	15	4	6
(13)	12	13	16	5	7
(14)	13	14	17	6	8
(15)	14	15	18	7	9
(16)	15	16	19	8	10
(17)	16	17	20	9	11
(18)	17	18	21	10	12
(19)	18	19	1	11	13
(20)	19	20	2	12	14
(21)	20	21	3	13	15

Plan 13.14 $t = 31, k = 6, r = 6, b = 31, \lambda = 1, E = 86$, Type I

Reps.

Block	I	II	III	IV	V	VI
(1)	31	1	3	8	12	18
(2)	1	2	4	9	13	19
(3)	2	3	5	10	14	20
(4)	3	4	6	11	15	21
(5)	4	5	7	12	16	22
(6)	5	6	8	13	17	23
(7)	6	7	9	14	18	24
(8)	7	8	10	15	19	25
(9)	8	9	11	16	20	26
(10)	9	10	12	17	21	27
(11)	10	11	13	18	22	28
(12)	11	12	14	19	23	29
(13)	12	13	15	20	24	30
(14)	13	14	16	21	25	31
(15)	14	15	17	22	26	1
(16)	15	16	18	23	27	2
(17)	16	17	19	24	28	3
(18)	17	18	20	25	29	4
(19)	18	19	21	26	30	5
(20)	19	20	22	27	31	6
(21)	20	21	23	28	1	7
(22)	21	22	24	29	2	8
(23)	22	23	25	30	3	9
(24)	23	24	26	31	4	10
(25)	24	25	27	1	5	11
(26)	25	26	28	2	6	12
(27)	26	27	29	3	7	13
(28)	27	28	30	4	8	14
(29)	28	29	31	5	9	15
(30)	29	30	1	6	10	16
(31)	30	31	2	7	11	17

Plan 13.15 $t = 37, k = 9, r = 9, b = 37, \lambda = 2, E = 91$, Type I

Reps.

Block	I	II	III	IV	V	VI	VII	VIII	IX
(1)	1	7	9	10	12	16	26	33	34
(2)	2	8	10	11	13	17	27	34	35
(3)	3	9	11	12	14	18	28	35	36
(4)	4	10	12	13	15	19	29	36	37
(5)	5	11	13	14	16	20	30	37	1
(6)	6	12	14	15	17	21	31	1	2
(7)	7	13	15	16	18	22	32	2	3
(8)	8	14	16	17	19	23	33	3	4
(9)	9	15	17	18	20	24	34	4	5
(10)	10	16	18	19	21	25	35	5	6
(11)	11	17	19	20	22	26	36	6	7
(12)	12	18	20	21	23	27	37	7	8
(13)	13	19	21	22	24	28	1	8	9
(14)	14	20	22	23	25	29	2	9	10
(15)	15	21	23	24	26	30	3	10	11
(16)	16	22	24	25	27	31	4	11	12
(17)	17	23	25	26	28	32	5	12	13
(18)	18	24	26	27	29	33	6	13	14
(19)	19	25	27	28	30	34	7	14	15
(20)	20	26	28	29	31	35	8	15	16
(21)	21	27	29	30	32	36	9	16	17
(22)	22	28	30	31	33	37	10	17	18
(23)	23	29	31	32	34	1	11	18	19
(24)	24	30	32	33	35	2	12	19	20
(25)	25	31	33	34	36	3	13	20	21
(26)	26	32	34	35	37	4	14	21	22
(27)	27	33	35	36	1	5	15	22	23
(28)	28	34	36	37	2	6	16	23	24
(29)	29	35	37	1	3	7	17	24	25
(30)	30	36	1	2	4	8	18	25	26
(31)	31	37	2	3	5	9	19	26	27
(32)	32	1	3	4	6	10	20	27	28
(33)	33	2	4	5	7	11	21	28	29
(34)	34	3	5	6	8	12	22	29	30
(35)	35	4	6	7	9	13	23	30	31
(36)	36	5	7	8	10	14	24	31	32
(37)	37	6	8	9	11	15	25	32	33

Plan 13.16 $t = 3, k = 5, r = 5, b = 3, \lambda = 5, E = 96$, Type III

Reps.

Block	I	II	III	IV	V
(1)	1	2	3	2	3
(2)	2	3	1	1	2
(3)	3	1	2	3	1

Plan 13.17 $t = 3, k = 7, r = 7, b = 3, \lambda = 7, E = 98$, Type IV

Reps.

Block	I	II	III	IV	V	VI	VII
(1)	1	2	3	1	2	3	1
(2)	2	3	1	3	1	2	2
(3)	3	1	2	2	3	1	3

Plan 13.18 $t = 3, k = 8, r = 8, b = 3, \lambda = 8, E = 98$, Type III

Reps.

Block	I	II	III	IV	V	VI	VII	VIII
(1)	1	2	3	1	2	3	1	2
(2)	2	3	1	3	1	2	2	3
(3)	3	1	2	2	3	1	3	1

Plan 13.19 $t = 3, k = 10, r = 10, b = 3, \lambda = 10, E = 99$, Type IV

Reps.

Block	I	II	III	IV	V	VI	VII	VIII	IX	X
(1)	1	2	3	1	2	3	1	2	3	1
(2)	2	3	1	3	1	2	2	3	1	3
(3)	3	1	2	2	3	1	3	1	2	2

Plan 13.20 $t = 4, k = 5, r = 5, b = 4, \lambda = 5, E = 96$, Type IV

Reps.

Block	I	II	III	IV	V
(1)	1	2	3	4	1
(2)	2	3	4	1	4
(3)	3	4	1	2	3
(4)	4	1	2	3	2

Plan 13.21 $t = 4, k = 7, r = 7, b = 4, \lambda = 7, E = 98$, Type III

Reps.							
Block I	II	III	IV	V	VI	VII	
(1)	1	2	3	4	1	2	3
(2)	2	1	4	3	3	4	1
(3)	3	4	1	2	4	3	2
(4)	4	3	2	1	2	1	4

Plan 13.22 $t = 4, k = 9, r = 9, b = 4, \lambda = 9, E = 99$, Type IV

Reps.								
Block I	II	III	IV	V	VI	VII	VIII	IX
(1)	1	2	3	4	1	2	3	4
(2)	2	1	4	3	3	4	1	2
(3)	3	4	1	2	4	3	2	1
(4)	4	3	2	1	2	1	4	3

Plan 13.23 $t = 5, k = 6, r = 6, b = 5, \lambda = 6, E = 97$, Type IV

Reps.						
Block I	II	III	IV	V	VI	
(1)	1	2	3	4	5	1
(2)	2	4	5	3	1	3
(3)	3	1	2	5	4	4
(4)	4	5	1	2	3	2
(5)	5	3	4	1	2	5

Plan 13.24 $t = 5, k = 9, r = 9, b = 5, \lambda = 9, E = 99$, Type III

Reps.								
Block I	II	III	IV	V	VI	VII	VIII	IX
(1)	1	2	3	4	5	1	2	3
(2)	2	3	4	5	1	4	5	1
(3)	3	4	5	1	2	2	3	4
(4)	4	5	1	2	3	5	1	2
(5)	5	1	2	3	4	3	4	5

Plan 13.25 $t = 6, k = 7, r = 7, b = 6, \lambda = 7, E = 98$, Type IV

Reps.							
Block	I	II	III	IV	V	VI	VII
(1)	1	2	3	4	5	6	1
(2)	2	3	6	1	4	5	3
(3)	3	6	2	5	1	4	5
(4)	4	5	1	2	6	3	4
(5)	5	1	4	6	3	2	6
(6)	6	4	5	3	2	1	2

Plan 13.26 $t = 7, k = 8, r = 8, b = 7, \lambda = 8, E = 98$, Type IV

Reps.								
Block	I	II	III	IV	V	VI	VII	VIII
(1)	1	2	3	4	5	6	7	1
(2)	2	5	1	7	6	4	3	3
(3)	3	6	7	2	4	1	5	7
(4)	4	7	5	6	3	2	1	6
(5)	5	4	2	3	1	7	6	5
(6)	6	3	4	1	7	5	2	4
(7)	7	1	6	5	2	3	4	2

CHAPTER 14

ANALYSIS OF THE RESULTS OF A SERIES OF EXPERIMENTS

14.1 Initial Steps in the Analysis

14.11 Introduction. In a program of research it is quite common to repeat the same experiment at a number of different places, on a number of different occasions. There may be several reasons for this. Sometimes the object of the research is to produce recommendations which are to apply to a population that is extensive either in space or in time or in both. Thus in agricultural field experimentation, many projects are undertaken in the hope that their results can be applied in practical farming. The conclusions drawn from such research, if they are to be of use, must be valid for at least several seasons in the future and over a reasonably large area of farm land. It has been found that the effectiveness of the common plant nutrients, of different varieties of a crop, and of different cultivation practices usually varies from field to field and, even more markedly, from season to season. A single experiment, however well conducted, supplies information about only one place and one season. Consequently such experiments are carried out at several different places in the area for which recommendations are wanted, and are repeated for a number of seasons.

In other cases we may be interested, not in making inferences about some specific population, but in studying the influence of external conditions on some measurement or on the responses to treatments. For example, is the vitamin A content of a vegetable affected by the climate in which it is grown? Do the relative durabilities of two types of road surface depend on weather or topography or traffic load? Repetition of such experiments in different places is necessary in order to have variations in the external factors that are under investigation.

A third example is supplied by collaborative experiments on biological assay by a group of laboratories. Here the objects may be to obtain information about the accuracy with which the potency of a drug can be estimated and to discover whether different laboratories reach the same conclusions about the relative potencies of different drugs. Bliss (14.1)

gives an interesting account of a series of experiments of this type, where sixteen laboratories estimated the potency of preparations of digitalis by injection into cats.

The appropriate statistical analysis for the data from a series of experiments will of course vary with the object of the research. Nevertheless, the preliminary stages of the analysis tend to be the same in all cases. In this chapter an introductory account is given of these initial procedures, which present difficulties that are not always appreciated. The first step, which is to analyze and interpret the data from each individual experiment, will be assumed to have been completed.

When we begin to combine the data, the first point of interest is to examine whether the differences among treatments are the same in all experiments. This question is likely to be relevant whatever the purpose of the experiments. In experiments designed to lead to the recommendation of a "best" treatment for some operation, we wish to know whether there is a consistent superiority of certain treatments, or whether on the other hand we may have to consider recommending different treatments for different circumstances. In the other two types of experimentation mentioned above, the test would indicate whether the responses to treatments have varied with the external conditions of the experiment, or whether the different laboratories agreed in their estimates of relative potency.

Secondly, it is often, though not always, desirable to estimate and compare the average effects of treatments over the whole series of experiments.

14.12 Numerical Example. The data come from a group of 6 experiments on Irish potatoes conducted in two counties of North Carolina in 1945 and 1946. The results have been described by Nelson and Hawkins (14.2), and form part of a study of the responses to applications of superphosphate on soils of varying degree of fertility. In 1945, on each of the 6 sites, the amount of readily soluble phosphorus in the soil itself was estimated from soil samples by the modified Truog method. These amounts varied from 48 to 850 lb. P_2O_5 per acre. The treatments comprised 5 different levels of application, 0, 40, 80, 120, and 160 lb. P_2O_5 per acre.

These experiments are an example of the second type discussed above, in that their object was to find out the extent to which the responses to treatments were influenced by the condition of the soil. The soils were not intended to be a representative sample of the soils in the counties, but were chosen so as to give a wide range in fertility.

Potatoes were grown on the plots in 1945 and 1946, the treatments being applied in both years, with in addition substantial applications of nitrogen and potash on all plots so that these nutrients would not be deficient. The data to be analyzed are the 1946 results. The mean yields and error m.s. per plot are shown in table 14.1. In five of the six experi-

TABLE 14.1 TREATMENT YIELDS (100 LB. PER ACRE) OF IRISH POTATOES, 1946

Pounds P_2O_5 applied per acre	Experiment number					
	1	2	3	4	5	6
	Amount of readily soluble phosphorus (pounds P_2O_5) in soil, 1945					
	48	310	410	710	790	850
0	114	142	180	170	130	170
40	206	176	188	171	140	178
80	231	201	207	188	150	187
120	237	205	208	185	152	188
160	252	217	217	189	156	189
s_e^2 *	106	83	158	43	204	52
d.f.	21	12	12	12	12	12
r †	4	5	5	5	5	5

* Error m.s. per plot.

† Number of replications.

ments the F -ratios for treatments against error were significant. One feature of the results in table 14.1 is that experiment 1, which has the lowest amount of available phosphorus in the soil according to the soil tests, actually gave the highest total yield in 1946. It should be remembered that the soil tests were made in 1945, and that some residual effects of the 1945 applications may have persisted. Whatever the reason, this soil was apparently very responsive to phosphorus in 1946; the plots without phosphorus do have the lowest yield of all places.

It is possible to compute from table 14.1 a combined analysis of variance for all six experiments. As will be seen later, there are limitations to the use of such an analysis. For the present we will ignore any complexities that may arise. Further, in order to discuss only the simplest case at first, we will omit experiment 1, which differs in size and structure from the other experiments. This omission is of course inappropriate from the agronomic point of view, but the data are being used to throw light on the general procedure in analysis rather than on the agronomic questions involved.

14.13 Preliminary Combined Analysis When all Experiments Have the Same Design. Formally, the analysis subdivides into the following components.

Places
Treatments
Treatments \times places
Pooled experimental error

In addition, in a complete analysis there are terms representing the differences among rows and columns in the individual experiments. Since these are not relevant to the interpretation, they are not included.

The analysis can be computed from either treatment means or totals; the totals will be used here and are shown in table 14.2.

TABLE 14.2 TREATMENT TOTALS FOR EXPERIMENTS 2 TO 6 (IN 100 LB.)

Pounds P ₂ O ₅	Experiment					Total
	2	3	4	5	6	
0	710	900	850	650	850	3,960
40	880	940	855	700	890	4,265
80	1,005	1,035	940	750	935	4,665
120	1,025	1,040	925	760	940	4,690
160	1,085	1,085	945	780	945	4,840
Total	4,705	5,000	4,515	3,640	4,560	22,420

The first 3 components are computed in the same way as for a single randomized blocks experiment, except that all sums of squares are divided by an extra 5 in order to reduce them to a single-plot basis. Thus we have

$$\text{Places: } \frac{(4705)^2 + \dots + (4560)^2}{25} - \frac{(22,420)^2}{125} = 41,367$$

$$\text{Treatments: } \frac{(3960)^2 + \dots + (4840)^2}{25} - \frac{(22,420)^2}{125} = 20,979$$

To obtain the treatments \times places s.s. we calculate the total s.s. for table 14.2, which comes to 69,199. Then

$$\text{Treatments } \times \text{ places: } 69,199 - 41,367 - 20,979 = 6853$$

Since all experiments have the same number of error degrees of freedom, the pooled error m.s. may be found as the simple average of the five s^2 values in table 14.1. This comes to 108, as given in table 14.3.

The null hypothesis that the treatment differences are the same at all places (i.e., that there are no treatment \times place interactions) is tested

by the F -ratio 428/108, or 3.96, with 16 and 60 d.f., respectively. Since the 5% level is 1.81, the ratio is definitely significant.

TABLE 14.3 COMBINED ANALYSIS OF VARIANCE (ON A SINGLE-PLOT BASIS)

Source of variation	d.f.	s.s.	m.s.
Places	4	41,367	
Treatments	4	20,979	5,245
Treatments \times places	16	6,853	428
Pooled error	60	6,480	108

A test of the average responses to treatments taken over all five places is of minor interest in this example, because the places do not constitute a random sample from a population about which we wish to make inferences. For purposes of illustration, however, we will assume that the places *were* selected as a random sample of fields on which potatoes might be grown commercially.

In the F -test of the average effects of treatments there are two possible candidates for the denominator of F —the mean square for the treatments \times places interactions (428) or the pooled error m.s. (108). Sometimes this competition does not arise, because our knowledge of the data and the F -test of the interactions both indicate that there is no reason to suppose real interactions to be present. In that event the mean squares for interactions and error may be pooled to form a single denominator for the F -test of treatments. But frequently the experimental conditions are such that we expect interactions to be present. This is so in the example, where previous work in a number of countries has shown a relation between the response to phosphorus and the amount found in the soil by a soil test. Consequently, even if the interactions m.s. had not proved significant, we might have been unwilling to assume that there were no interactions.

In discussing the two F -ratios it is helpful to examine the mathematical model on which the combined analysis is based. If x_{ij} is the observed mean for the j th treatment at the i th place, we postulate that

$$x_{ij} = \mu + \pi_i + \tau_j + \mu_{ij} + \bar{\epsilon}_{ij} \quad (14.1)$$

where π_i , τ_j represent the effects of the place and the treatment, respectively, μ_{ij} that of the treatment \times place interaction, and $\bar{\epsilon}_{ij}$ that of the experimental error. $\bar{\epsilon}_{ij}$ is the average of the errors on the r plots that receive the treatment at that place.

With this model we can study the nature of the mean squares in the analysis of variance. If the experimental errors on individual plots have a variance σ_e^2 , and if the interaction terms μ_{ij} may be considered to have

a variance σ_μ^2 , the average values of the mean squares work out as follows.

TABLE 14.4 EXPECTED VALUES OF MEAN SQUARES

Source of variation	Expected value of mean square
Treatments	$\sigma_e^2 + r\sigma_\mu^2 + \frac{rp}{(t-1)} \sum (\tau_j - \bar{\tau})^2$
Treatments \times places	$\sigma_e^2 + r\sigma_\mu^2$
Pooled experimental error	σ_e^2

The symbols r , p , and t stand respectively for the numbers of replications, places, and treatments.

It will be noted that the treatments m.s. is influenced by three components—the experimental error variance, the variance of treatment \times place interactions, and the variance among the true treatment means τ_j . The pooled experimental error takes account of only the first of these components. Hence the treatments m.s. may be statistically significant, as compared with the pooled error m.s., either because there are real differences among the τ 's or because treatment \times place interactions are present. This F -test is informative only when we are indifferent as to whether a significant treatments m.s. was due to real treatment differences or to interactions. This situation is rare in practice. In most cases, on the contrary, having established that interactions are present, or at least that it is not safe to assume them absent, we wish to know whether *in addition* there are consistent differences among the effects of treatments. As table 14.4 shows, the appropriate denominator of F for this test is the interactions m.s. This contains both the interaction and error components of variation in exactly the same way as they enter into the treatments m.s. The F -ratio, 5245/428, or 12.26, with 4 and 16 d.f., is significant at the 1% level.

The conclusions from this initial analysis are: (i) there are real differences in response that are consistent from place to place, and (ii) there are real variations in responsiveness from place to place. It need not be stressed that these statements do not constitute a competent summary of the results. They merely indicate what should be examined next. In the present case we would consider which treatments have proved consistently superior, and, if possible, why this is so. Similarly, we should investigate the nature of the interactions and try to find a rational explanation for them.

14.2 Criticisms of the Preliminary Analysis

14.21 Heterogeneity of the Interaction Variance. The previous analysis is open to several criticisms. Although these criticisms are not valid

for all series of experiments, our experience is that it is well to be on one's guard against them. They deal essentially with the assumptions on which the combined analysis was based.

These assumptions have been presented in section 14.13. Specifically, we postulate that the model (14.1) holds. Further, the experimental errors e_{ij} of individual observations are assumed to be normally and independently distributed with the same variance σ_e^2 . Finally, for the F -test of the treatments m.s. against the treatments \times places interactions, we require also the assumptions that the interaction terms μ_{ij} are normally and independently distributed, with zero population means and the same variance σ_μ^2 , and are independent of the e 's.

The first criticism is that some components of the treatments \times places s.s. may be much larger than others, or in mathematical terms that the "interaction" variance σ_μ^2 is not constant. This will happen if the effectiveness of some treatments varies greatly from place to place, while that of others varies less or not at all.

If the interaction m.s. is heterogeneous in this sense, the F -test of treatments against interactions is vitiated. The general effect is that the F value read from the tables is too low, i.e., that too many significant results are obtained. Some idea of the extent of the bias can be obtained in certain extreme cases. In the numerical example, if one comparison among the treatment means has a much larger interaction variance than any other comparisons, the F -ratio is distributed approximately as an F value with 1 and 4 d.f., respectively, instead of 4 and 16 d.f. The 5% significance level would be 7.71 instead of 3.01. More generally, if there are p places, F is distributed approximately with 1 and $(p - 1)$ d.f. This situation produces about the greatest distortion in F that is likely to arise, so that 7.71 could be regarded as an upper limit to the significance level of F .

The exact distribution of F can be worked out, but is not available in a form adapted for practical use. Even so, in many instances, uncertainty about the correct significance level for F does not preclude us from drawing conclusions from the test. For, in the example above, if F turned out to be 1.38 we would be confident that it is *not* significant, since the 5% level is at least 3.01. Similarly, there seems little doubt that the F value actually obtained in the experiments, 12.26, *is* significant, since it lies well above 7.71. On the other hand, we would be uncertain about significance if F happened to lie between 3 and 7.

A better method of coping with this difficulty is to divide the treatment s.s. into a set of orthogonal components that will supply all or most of the information of interest to us. The interactions s.s. is partitioned in the same way so as to isolate the interactions of each component with places. By Bartlett's test for homogeneity of variance, reference (14.3),

we can then test whether σ_μ^2 is the same for all the components of the interactions s.s. If we decide that σ_μ^2 can be assumed constant, the difficulty vanishes. If σ_μ^2 is not constant, it is valid to test any component of the treatments s.s. against *its own* interaction with places. Apart from the extra computations, the only drawback to this procedure is that the degrees of freedom in the denominator of F are reduced.

These remarks will be illustrated by the example. In these experiments the principal point of interest is to see whether the average response to a given amount of phosphorus decreases when the amount presumed to be in the soil increases. It may also be worth while to examine whether the rate of decline in response with the higher levels of dressing changes with the nature of the soil. These questions can be considered conveniently by fitting at each place a parabolic regression of the yield on the amount of dressing. Although a parabolic regression would scarcely be regarded as the true form of the response curve, it appears to fit the data well.

Since the dressings increase by equal amounts, the orthogonal polynomials given by Fisher and Yates (14.4) are suitable. The calculations are shown below for experiment 3. The linear term is proportional to

Amount of P_2O_5	Total yield	Multipliers for	
		Linear term	Quadratic term
0	900	-2	2
40	940	-1	-1
80	1035	0	-2
120	1040	1	-1
160	1085	2	2
Sum of products		470	-80
Divisor for square		50	70

the average increase in yield per 40 lb. of P_2O_5 : the quadratic term may be interpreted as measuring the rate of decline in response with increased dressings. The values found for the two terms are as follows.

	Experiment					Total
	2	3	4	5	6	
L	895	470	260	320	240	2185
Q	-325	-80	-70	-100	-110	-685

The contributions to the treatments s.s. are

$$\text{Linear: } \frac{(2185)^2}{250} = 19,097; \quad \text{quadratic: } \frac{(685)^2}{350} = 1341$$

In the treatments \times places s.s. we have

$$L \times \text{places: } \frac{(895)^2 + \dots + (240)^2}{50} - 19,097 = 5893$$

$$Q \times \text{places: } \frac{(325)^2 + \dots + (110)^2}{70} - 1341 = 645$$

In table 14.5 the three interaction mean squares are in the same order of size as the corresponding treatment mean squares. This result is typical, in that large effects tend to have large interactions. It is obvious on inspection that the components of the interaction m.s. cannot be regarded as homogeneous. Accordingly, we test the linear component of treatments against the mean square 1473 instead of against the complete treatments \times places m.s. of 428 in table 14.3. This makes a substantial difference to the F -ratio, though it remains significant. The quadratic \times places m.s. (161) is not significantly above the error m.s. (108), but it seems prudent to use it as the denominator in the F -test of the quadratic term in treatments. The remainder of the interactions shows no sign of significance, and could be pooled with error for an F -test of the deviations from regression.

TABLE 14.5 SUBDIVISION OF THE ANALYSIS OF VARIANCE

Source of variation	d.f.	s.s.	m.s.
Treatments			
Linear	1	19,097	19,097
Quadratic	1	1,341	1,341
Deviations from regression	2	541	270
Treatments \times places			
Linear \times places	4	5,893	1,473
Quadratic \times places	4	645	161
Deviations \times places	8	315	39
Pooled error	60	6,480	108

14.22 Heterogeneity of the Experimental Error Variances. A second criticism concerns the assumption that the experimental error variances σ_e^2 are the same in all experiments. In general this assumption will hold only if all experiments have been conducted in the same way, with the same amount of control over environmental conditions and with experimental material of the same variability. In experiments with crops or animals this degree of uniformity is seldom attainable, because the natural variability among pieces of land or among animals at one place differs from that at other places. Hence in this and many other types of cooperative experimentation we expect *a priori* that experimental error variances change from place to place.

When there is doubt about this point, Bartlett's test of homogeneity of variances can be applied to the error m.s., s^2 , in the experiments. In the example (table 14.1) the s^2 values varied from 43 in experiment 4 to 204 in experiment 5, and the test of homogeneity shows that σ_e^2 cannot be assumed constant.

Variation in σ_e^2 invalidates the F -test of the interactions m.s. against the pooled error m.s. Although the effect on the significance level of F is not known exactly, it operates so that use of the tabular F produces too many significant results. As in the previous section, the extent of the distortion in F can be seen in extreme cases. If one experiment has a much higher error variance than any of the others, F will be distributed approximately as the tabular F with $(t - 1)$ and n' degrees of freedom, where t is the number of treatments and n' is the number of error degrees of freedom in the experiment with the high error variance. In the example, if one of the experiments with 12 error d.f. happened to be much less accurate than the other experiments, F would have 4 and 12 d.f. instead of 16 and 60 as used in the test in section 14.13. The 5% level would be 3.26 instead of 1.81. Since this case appears to be the most unfavorable that would occur, we may conclude that the true significance level of F lies somewhere between these values.

If the observed F in our data falls outside these limits, a definite conclusion can be reached from the test without further knowledge of the exact significance level of F . This happens in the example, where the observed F -ratio for the interactions m.s. is 3.96. However, even if the exact significance level were known, the test can be criticized because the F -ratio is no longer the most sensitive test criterion. Studies have shown that, with the amount of variation in σ_e^2 that appears typical of agricultural experimentation, this loss of sensitivity might be equivalent to discarding 10 to 20% of the data. An approximate test that avoids some of the loss in sensitivity will be presented in section 14.4.

Thus far we have been discussing how the F -test of the interactions is affected by heterogeneity in the error variances. The F -test of treatments may now be considered. If the interactions are regarded as negligible, so that the denominator of F is the pooled m.s. for interactions and error, the F -test of treatments is invalidated in about the same way as that of interactions. On the other hand, when interactions are present, and especially when they are large, the F -test of the treatments m.s. against the interactions m.s. is much less disturbed. The reason may be clearer if we consider the test of the linear component of treatments in table 14.5. The suggested denominator for this test was the linear \times places component, 1473. This is nearly 14 times the pooled error m.s. It appears, therefore, that heterogeneity in the error variances

can influence only a small part of the interaction m.s., so that its effect is, as it were, greatly reduced. These remarks apply with much less force to the quadratic \times places m.s., to which, taking the data at their face value, the pooled error contributes more than half.

14.23 Summary. When all experiments are identical in structure, a combined analysis of variance can be computed relatively easily. Application of the ordinary tests of significance to this analysis is frequently open to question because of heterogeneity in the error and interaction variances. Despite this, it is advisable to draw preliminary conclusions as far as possible from this analysis, at least in the present state of our knowledge, because tests that are fully efficient and theoretically sound have not yet been discovered, and the approximate tests that have been devised to meet the criticisms are more laborious.

When the interactions are sizable, there is usually little difficulty in interpreting the combined analysis. For, if the F -ratio for interactions is large, we may be confident that it is statistically significant in spite of some uncertainty about the exact significance level of F , and the criticism that the F -test is not fully sensitive carries less weight if F establishes significance. Further, as we have seen, the F -test of the treatments m.s. against the interactions m.s. is little affected by inequality in the error variances when the interactions are large. The chief point to remember is that the interactions m.s. may itself be heterogeneous. Subdivision of the treatments and interactions m.s. according to the treatment comparisons that are of greatest importance is often useful.

Another situation that presents little difficulty occurs when the average differences among treatments are substantial, yet interactions are negligible. In this case it may be found that neither the F value for interactions nor that for treatments is close to the significance level. The cases that leave us in doubt are those where the F values are just slightly above the tabular significance levels.

14.3 Experiments of Unequal Size

14.31 Numerical Example. Provided that the same set of treatments appears in all experiments, a combined analysis can usually be constructed even when the experiments differ in size and structure. Some issues arise, however, that are not encountered when all experiments are identical in design. The problems will be illustrated by the inclusion of the first experiment in the previous example. This was arranged in randomized blocks with 4 replications, whereas all other experiments were in

5 × 5 latin squares*. The treatment totals and numbers of replications are shown in table 14.6.

TABLE 14.6 TREATMENT TOTALS (in 100 lb.)

Pounds P ₂ O ₅	Experiment						Total	No. of plots
	1	2	3	4	5	6		
0	456	710	900	850	650	850	4,416	29
40	824	880	940	855	700	890	5,089	29
80	924	1,005	1,035	940	750	935	5,589	29
120	948	1,025	1,040	925	760	940	5,638	29
160	1,008	1,085	1,085	945	780	945	5,848	29
Total	4,160	4,705	5,000	4,515	3,640	4,560	26,580	145
No. of plots	20	25	25	25	25	25		

A combined analysis can be obtained by following the standard procedure for data based on unequal numbers. The square of any quantity is divided by the number of replications involved. The sums of squares are given below.

$$\text{Total: } \frac{(456)^2 + \dots + (1008)^2}{4} + \frac{(710)^2 + \dots + (945)^2}{5} - \frac{(26,580)^2}{145} = 131,925$$

$$\text{Places: } \frac{(4160)^2}{20} + \frac{(4705)^2 + \dots + (4560)^2}{25} - \frac{(26,580)^2}{145} = 55,509$$

$$\text{Treatments: } \frac{(4416)^2 + \dots + (5848)^2}{29} - \frac{(26,580)^2}{145} = 45,613$$

$$\text{Treatments} \times \text{places: } 131,925 - 55,509 - 45,613 = 30,803$$

The pooled error m.s. requires a little thought. If all experiments have the same true error variance σ_e^2 , the best procedure is to weight each error m.s., s_i^2 , by the number of degrees of freedom n_i . But if the experiments vary in accuracy, this weighted mean is a biased estimate of the component of error variance that enters into the interaction and treatments m.s. The correct component is $\sum r_i \sigma_i^2 / \sum r_i$ instead of $\sum n_i \sigma_i^2 / \sum n_i$. Consequently, unless we are confident that σ_e^2 is constant, it is best to weight the s_i^2 values by the numbers of replications. This gives

$$\bar{s}_e^2 = \frac{4(106) + 5(83 + 158 + \dots + 52)}{29} = 108$$

* Since experiment 1 contained additional treatments (not discussed here), it provided 21 error d.f.

The F -test of the interactions m.s. against the error m.s. is carried out in the same way as with experiments of equal size, and is subject to the criticisms previously discussed. The F -test of the treatments m.s. against the interactions m.s. encounters a new difficulty that is due to

TABLE 14.7 PRELIMINARY ANALYSIS OF VARIANCE FOR EXPERIMENTS OF UNEQUAL SIZE

Source of variation	d.f.	S.S.	m.s.
Places	5	55,509	11,102
Treatments	4	45,613	11,403
Treatments \times places	20	30,803	1,540
Pooled error m.s.	81		108

the unequal numbers of replications and is present even if all the assumptions required for the analysis of variance are satisfied. Under these assumptions, the expectations of the principal mean squares in table 14.7 are as given in table 14.8.

TABLE 14.8 EXPECTED VALUES OF MEAN SQUARES (WITH UNEQUAL NUMBERS OF REPLICATIONS)

Source of variation	Expected value of mean square
Treatments	$\sigma_e^2 + \bar{r}_1\sigma_\mu^2 + (\sum r_i) \frac{\sum (\tau_j - \bar{\tau})^2}{(t-1)}$
Treatments \times places	$\sigma_e^2 + \bar{r}_2\sigma_\mu^2$
Pooled error	σ_e^2

where

$$\bar{r}_1 = \frac{\sum r_i^2}{\sum r_i} = \frac{141}{29} = 4.862$$

$$\bar{r}_2 = \frac{1}{(p-1)} (\sum r_i - \bar{r}_1) = \frac{1}{5} (29 - 4.862) = 4.828$$

The complicating factor is that the coefficient of σ_μ^2 in the expected treatments m.s. is not the same as that in the expected treatments \times places m.s.; in fact, it is always larger. The difference is very small in this example, as it is whenever the experiments do not vary much in numbers of replications. In addition, and for the same reason, the interactions m.s. is not distributed as a multiple of chi-square; that is, it does not have the type of distribution required for the validity of the F -test. Both factors tend to make the F -test give too many significant results.

A crude method which at least prevents us from being led too far astray is to adjust the F -ratio so as to remove the upward bias in expectation. Let the expected values of the three mean squares be de-

noted by θ_t , θ_{tp} , and θ_e , respectively. Then, if the null hypothesis is true (all τ_j equal), it is seen from table 14.8 that the three expected values are connected by the equation

$$\theta_t + (k - 1)\theta_e = k\theta_{tp}$$

where $k = \bar{r}_1/\bar{r}_2$. In other words, the test that we seek is a test that this relation holds. This suggests that F might be computed as

$$F' = \frac{s_t^2 + (k - 1)\bar{s}_e^2}{ks_{tp}^2} = \frac{11,403 + 0.007(108)}{(1.007)(1540)} = 7.35$$

as compared with the original F -ratio of 11,403/1540, or 7.40. The significance level of F' will also be altered slightly from that in the table.

When interactions are likely to be large, and the chief interest centers in the test of the treatments m.s. against the interactions m.s., an alternative approach has much to commend it. This is to compute the analysis from the *unweighted* treatment means given in table 14.1, paying no attention to the numbers of replications. With this type of analysis the bias disappears, the expectations of the mean squares being shown in table 14.9. It will be noted that σ_e^2 and σ_μ^2 carry the same coefficients

TABLE 14.9 EXPECTED VALUES OF MEAN SQUARES IN AN UNWEIGHTED ANALYSIS

Source of variation	Expected value of mean square
Treatments	$\frac{\sigma_e^2}{\bar{r}_h} + \sigma_\mu^2 + p \frac{\sum (r_j - \bar{r})^2}{(t - 1)}$
Treatments \times places	$\frac{\sigma_e^2}{\bar{r}_h} + \sigma_\mu^2$

in both expectations. The divisor \bar{r}_h is the harmonic mean of the numbers of replications, given by $\bar{r}_h = p / \sum (1/r_i)$. Moreover, if σ_μ^2 is much larger than σ_e^2/\bar{r}_h , the interactions m.s. tends to be distributed as a multiple of chi-square, so that the conditions for the F -test are closely approximated.

It is suggested that the pooled error m.s. for insertion in an analysis of this type be calculated from the formula

$$\bar{s}_e^2 = \frac{1}{p} \left(\frac{s_1^2}{r_1} + \frac{s_2^2}{r_2} + \cdots + \frac{s_p^2}{r_p} \right)$$

By an extension of the results in table 14.9 to the case where the experiments have different error variances, this quantity is found to be an un-

biased estimate of the error component that enters into the mean squares for treatments and treatments \times places. For the data in table 14.1 we have

$$\bar{s}_e^2 = \frac{1}{6} \left(\frac{106}{4} + \frac{83 + 158 + \cdots + 52}{5} \right) = 22.4$$

TABLE 14.10 UNWEIGHTED ANALYSIS OF VARIANCE OF TREATMENT MEANS

Source of variation	d.f.	s.s.	m.s.
Places	5	11,692	2,338
Treatments	4	10,554	2,638
Treatments \times places	20	7,158	358
Pooled error	81		22.4

Table 14.10, being in terms of treatment means, is in different units from table 14.7 (p. 403). As a conversion factor, the harmonic mean \bar{r}_h , which in this case is 4.8, may be used to multiply the mean squares above for comparison with those in table 14.7, though this conversion is not needed for our purpose.

In a choice between a weighted and an unweighted analysis, the following are the relevant considerations. The statements below are strictly true only if all experiments have the same error variance per observation (or plot), though they should remain substantially true with a moderate variation in σ_i^2 . The weighted analysis is superior for the F -test of the interactions. It gives a more powerful test, and the F -ratio approximates the tabular distribution of F more closely. It is also superior for the F -test of treatments if interactions are non-existent or small, since in this case the bias in the test is negligible. The unweighted analysis is preferable for the F -test of treatments when the interactions are not negligible. As it happens, the results in the example are not too well in accord with these statements. For the interactions test, the F -ratio is 14.26 with the weighted analysis and 15.98 with the unweighted analysis. The explanation of the higher value with the unweighted analysis is probably that experiment 1, whose responses differed from those in the other experiments, had the smallest number of replications. The values of F in the test of treatments were practically identical, being 7.35 (after adjustment) in the weighted analysis and 7.37 in the unweighted analysis.

14.32 The Combined Analysis for a Series of Lattice Experiments. In the discussion of lattice experiments it was pointed out that if inter-block information is recovered, the F -test of treatments in an individual ex-

periment is not exact, because the relative weights attached to inter- and intra-block estimates are subject to sampling errors. For the same reason any combined analysis must also be approximative rather than exact.

The designs need not be identical at all places. With 25 treatments, for example, some experiments might be simple lattices in 4 replications, while others are lattice squares in 3 replications. Also, inter-block information need not be recovered in all analyses; at some places we might have used a randomized blocks analysis.

If the experiments have different numbers of replications, the previous section has indicated that there is a question whether to use a weighted or an unweighted analysis. The former is preferable when the chief purpose is to test the interactions with places, and when such interactions are negligible; the latter when interactions are sizable and the principal interest is in a test of the average effects of treatments. With lattice experiments as used in agriculture, we are often interested in both interactions and average effects. But as interactions are seldom absent, the unweighted analysis is frequently advisable, and has the virtue of being slightly simpler.

The weighted analysis will be described first. First form a two-way "treatments \times places" table similar to table 14.6. The entries in the table are the *adjusted* treatment totals, except in experiments analyzed by randomized blocks, where unadjusted totals are used. This table is analyzed into components for

Places
Treatments
Treatments \times places

If the numbers of replications r_i differ, the weighted analysis follows the same procedure as in table 14.6 (p. 402). The square of every marginal treatment total is divided by $\sum r_i$ in order to reduce the analysis of variance to a single-observation basis. The square of a place total, summed over all treatments, is divided by tr_i , where t is the number of treatments.

The only additional component is the pooled error m.s. In any experiment where inter-block information has been recovered, we use as the estimate of the error variance per observation (or plot) the *effective* error m.s. E'_t . This quantity is obtained by adjusting the intra-block m.s. upwards so as to allow for sampling errors in the block corrections, and its calculation is explained along with the analysis for each type of

lattice. For the triple lattice, for example, E'_t is $E_e \left[1 + \frac{rk\mu}{k+1} \right]$. If there are any experiments in which the intra-block analysis has been carried out, E'_t is given the value which it takes when E_t becomes very

large relative to E_i . For experiments analyzed by randomized blocks, E'_i is the ordinary error m.s.

On the assumption that the true effective variances are unlikely to be the same in all experiments, the pooled error m.s. is calculated as $\sum r_i E'_i / \sum r_i$. When r_i is constant, this reduces to the unweighted mean of the E'_i . Tests of significance are made as in the numerical example, and are subject to the same general criticisms. Since the larger lattice designs provide ample degrees of freedom for error, the pooled error may be relatively well determined even if some experiments are much more accurate than others, so that there is less uncertainty about the true significance levels of the F -ratios.

For an unweighted analysis with experiments of unequal size, construct a two-way table of adjusted treatment *means*. This is analyzed by the simple standard procedure. The comparable pooled error m.s. is taken as

$$\bar{s}_e^2 = \frac{1}{p} \left(\frac{E_1'}{r_1} + \frac{E_2'}{r_2} + \cdots + \frac{E_p'}{r_p} \right)$$

With balanced designs, the method given above is a natural extension of the technique used to obtain an approximate F -test of treatments in an individual experiment. With partially balanced designs the method is more crude than that used in single experiments, where a special supplementary calculation was made for the F -test. This calculation can be extended to apply to a combined analysis, but it is doubtful whether the elaboration is worth while.

14.4 A Test of the Treatments \times Places Interactions

In this section we give an approximate test of the treatments \times places interactions for series of experiments where there is considerable variation in the experimental error variances. It was suggested that the F -test, despite its imperfections, will often serve our purpose. The present test differs from the F -test in that it gives less weight to experiments which have a high error variance, and consequently may be expected to be more sensitive. It may be useful in cases where there is doubt about the verdict given by the F -test and in cases where the most efficient analysis is desired.

As before, let x_{ij} be the mean of the j th treatment, and σ_i^2 be the true error variance per observation in the i th experiment. We assume that all experiments have the same design, and let s_i^2 be the error m.s. per observation in the i th experiment, based on n degrees of freedom.

For *known* values of σ_i^2 the theory of least squares indicates that the

mean x_{ij} should receive a weight $W_i = r/\sigma_i^2$. In a weighted analysis of variance of the treatment means, the interaction s.s. is known to be distributed as chi-square with $(p-1)(t-1)$ degrees of freedom. This sum of squares appears to be the best test criterion available, unless we possess specific information about the type of interaction that may exist, in which event a more specialized criterion would be constructed.

In default of knowledge of the σ_i^2 , the natural step is to consider a weighted analysis with weights $w_i = r/s_i^2$. That is, we weight each mean inversely as its *estimated* variance, since we do not know its true variance. This is satisfactory, provided that the s_i^2 are good estimates of the σ_i^2 . Upon examination it appears that the s_i^2 should be based on at least 15 d.f. To illustrate the calculations, the test will be applied to the data from the last 5 experiments, though the number of degrees of freedom, 12, is slightly too low to use the test with full confidence.

TABLE 14.11 TREATMENT MEAN YIELDS (100 LB. PER ACRE) FOR A WEIGHTED ANALYSIS OF VARIANCE

Pounds P ₂ O ₅	Experiment					Weighted total $\sum w_i x_{ij} = T_j$
	2	3	4	5	6	
0	142	180	170	130	170	53.570
40	176	188	171	140	178	57.000
80	201	207	188	150	187	62.194
120	205	208	185	152	188	62.264
160	217	217	189	156	189	63.932
w_i	0.060	0.032	0.116	0.025	0.096	0.329 = W
Total (P_i)	941	1000	903	728	912	
$w_i P_i$	56.460	32.000	104.748	18.200	87.552	298.960 = G
s.s. (S_i)	180,655	200,946	163,431	106,440	166,618	

The arrangement of the data in table 14.11 should be noted. The treatment means and s_i^2 values come from table 14.1 (p. 393). The weights for individual entries in the table are placed in the row immediately below the entries; for example, $w_1 = 5/83 = 0.060$. The total of the weights is denoted by W .

1. Form the column totals P_i and the products $w_i P_i$. Form the weighted row totals. The corner total, $G = 298.960$, supplies a check on both the column and row totals.

2. The items in the analysis of variance are obtained as follows.

Correction term:

$$C = \frac{G^2}{tW} = \frac{(298.960)^2}{(5)(0.329)} = 54,332.57$$

Total: First compute and record in the table the sum of squares S_i (uncorrected) of the entries in each column. The total s.s. is then found as

$$\sum (w_i S_i) - C = (0.060)(180,655) + \cdots + (0.096)(166,618) - C = 551.33$$

Places: A column (place) total has variance $t\sigma_i^2/r$, and hence receives a weight w_i/t . The sum of squares is

$$\frac{1}{5} \sum (w_i P_i^2) - C = 230.09$$

This is found readily by using the products $w_i P_i$.

Treatments: Each treatment total has estimated weight W . The sum of squares is

$$\frac{\sum T_j^2}{W} - C = \frac{(53.570)^2 + \cdots + (63.932)^2}{0.329} - C = 229.57$$

Treatments \times places: Since this is the only component in which we are currently interested, it is unfortunate that it must be found by means of the others. As usual, it is given by

$$551.33 - 230.09 - 229.57 = 91.67 = I$$

The sum of squares I does not follow a chi-square distribution, being inflated by errors in the weights. It can be reduced to a quantity that is distributed approximately as chi-square.* We take

$$\chi^2 = \frac{(n-4)(n-2)}{n(n+t-3)} I, \quad \text{with } \frac{(p-1)(t-1)(n-4)}{(n+t-3)} \text{ d.f.}$$

In this case

$$\chi^2 = \frac{(8)(10)}{(12)(14)} (91.67) = 43.65, \quad \text{with } \frac{(16)(8)}{(14)} = 9.14 \text{ d.f.}$$

In this approximation the degrees of freedom ascribed to chi-square are not integral. Significance levels are obtained by linear interpolation in the tables of chi-square. Since the 1% levels of chi-square are 21.67 and 23.21 for 9 and 10 d.f., respectively, I is obviously significant.

The test can also be applied to any component of the interactions s.s. As an example, we will test the interaction of the quadratic component of the regression on amount of phosphorus. This was previously tested by the F -test in table 14.5, and found non-significant. The values given

* This test is an extension and modification of a test previously given by Cochran (14.6).

for the quadratic components in section 14.21 were derived from treatment totals; to obtain corresponding values from means, we divide by 5.

	Experiment					
	2	3	4	5	6	Total
Q_i	-65	-16	-14	-20	-22	
$w_i Q_i$	-3.900	-0.512	-1.624	-0.500	-2.112	-8.648

Each quantity has variance $14 \sigma_i^2/r$, and receives a weight $w_i/14$. The sum of squares is

$$\frac{1}{14} \left(\sum w_i Q_i^2 - \frac{(\sum w_i Q_i)^2}{W} \right) = \frac{1}{14} (340.89 - 227.32) = 8.11 = I_q$$

The quantity I_q is the sum of squares for the interaction of a single treatment comparison with all p places, and would normally have $(p-1)$ degrees of freedom. To present the general formula for conversion to chi-square, suppose that we have computed the sum of squares for the interaction of t_1 treatment comparisons with p_1 of the places, so that there would normally be $t_1(p_1-1)$ degrees of freedom. Then

$$\chi^2 = \frac{(n-4)(n-2)}{n(n+t_1-2)} I_q, \quad \text{with } \frac{t_1(p_1-1)(n-4)}{(n+t_1-2)} \text{ d.f.}$$

This formula is in accord with that used for the whole $(t-1)(p-1)$ degrees of freedom in the interactions. Since the t treatments provide $(t-1)$ independent treatment comparisons, we would take $t_1 = (t-1)$ in applying the formula to the complete interactions s.s. For $t_1 = 1$, $p = 5$,

$$\chi^2 = \frac{(8)(10)}{(12)(11)} (8.11) = 4.92, \quad \text{with } \frac{(4)(8)}{(11)} = 2.91 \text{ d.f.}$$

The tables show that this value has a probability between 0.2 and 0.1, since by interpolation the 20% level for 2.91 d.f. is 4.51, the 10% level 6.10. The earlier F -test gave a probability slightly over 0.2, so that the two tests are in close agreement.

If the experiments differ in size, the weight w_i becomes r_i/s_i^2 . The numbers of error degrees of freedom will also vary. This necessitates a more elaborate conversion formula: as a rough approximation the average number of error degrees of freedom per experiment is used in place of n .

The test is not recommended for low values of n , because the s_i^2 are

relatively poor estimates of the σ_i^2 , and frequently one or two experiments receive such high weights that they dominate the analysis. Appropriate methods are presented by Yates and Cochran (14.5) and Cochran (14.6).

14.5 Repetitions in Both Space and Time

As has been mentioned, agricultural field experiments are often repeated both at a number of places and for a number of years. We will give only an introduction to the simplest case, in which experiments have been carried out at each of p places for y years. At any place a new site is chosen each year for the experiment, and a new randomization employed, so that the data from successive years may be regarded as independent. The experimental arrangement need not be uniform throughout the whole series of experiments: in fact, a change in design between seasons is not uncommon.

The mathematical representation of the mean of the j th treatment at the i th place in the k th year is now more lengthy. In addition to the symbols previously used for place and treatment effects, γ_k will denote the effect of the year. Interactions are denoted by multiple symbols: thus $(\pi\tau)_{ij}$ is the contribution of the place \times treatment interaction in this experiment. With this notation

$$x_{ijk} = \mu + \pi_i + \tau_j + \gamma_k + (\pi\tau)_{ij} + (\pi\gamma)_{ik} + (\tau\gamma)_{jk} + (\pi\tau\gamma)_{ijk} + \bar{e}_{ijk}$$

Note that there are treatment \times place, treatment \times year, and treatment \times place \times year interactions.

In experimental programs of this type, it is usually hoped that the places and years constitute a representative sample of the population of places and years to which the results will be applied. There are obvious practical difficulties in choosing places and years that can be confidently asserted to be such a representative sample, and sometimes little effort is made to ensure that this will be so. The hard fact is that any statistical inferences drawn from an analysis of the data will apply only to the population (if one exists) of which the experiments are a random sample. If this population is vague and unreal, the analysis is likely to be a waste of time, at least from the strictly practical point of view.

It seems appropriate to regard all the quantities in the equation as random variables except the general mean μ and the true effects τ_i of the treatments, because if we could tabulate all the values of say $(\pi\tau)_{ij}$ in the population, they would follow some frequency distribution from which the values in our data are a sample. For the full application of the analysis of variance we require the assumptions that the experi-

mental errors and all interactions of treatments are normally and independently distributed, with variances that are denoted by use of the letters that enter into the interaction.

An unweighted analysis of variance of the treatment means is obtained by the standard procedure for factorial experiments. The important parts of his analysis are sketched in table 14.12.

TABLE 14.12 ANALYSIS OF VARIANCE OF TREATMENT MEANS WITH TIME AND PLACE VARIATIONS

Source of variation	d.f.	Expectation of mean square
Treatments	$(t - 1)$	$\sigma_e^2 + \sigma_{tpy}^2 + p\sigma_{ty}^2 + y\sigma_{tp}^2 + py \frac{\sum (\tau_j - \bar{\tau})^2}{(t - 1)}$
Treatments × places	$(t - 1)(p - 1)$	$\sigma_e^2 + \sigma_{tpy}^2 + y\sigma_{tp}^2$
Treatments × years	$(t - 1)(y - 1)$	$\sigma_e^2 + \sigma_{tpy}^2 + p\sigma_{ty}^2$
Treatments × places × years	$(t - 1)(p - 1)(y - 1)$	$\sigma_e^2 + \sigma_{tpy}^2$
Pooled error		σ_e^2

In the most general case, σ_e^2 is the average value of σ_{ik}^2/r_{ik} , where σ_{ik}^2 is the error variance in the individual experiment at the i th place in the k th year, and r_{ik} is the number of replications in that experiment. Consequently, the pooled error should be estimated as the average of σ_{ik}^2/r_{ik} .

The method to be followed in testing the significance of the successive terms is made clear by the expected values of the mean square. The treatments × places × years interactions are tested against the pooled error. Both treatments × years and treatments × places are tested against treatments × places × years.

The situation with regard to the test for treatments is interesting. It is evident that no other mean square in the analysis is suitable as a denominator of F , unless either the treatments × places or the treatments × years interactions appear to be negligible, so that the corresponding variance is assumed to be zero. If this happens, the other interaction m.s. is an appropriate denominator for F (subject to some uncertainty in case the assumption should not be correct).

When both two-factor interactions are present, the hypothesis that all τ_j are zero is equivalent to the hypothesis that

$$\theta_t + \theta_{tpy} = \theta_{tp} + \theta_{ty}$$

where the θ 's stand for the expected mean squares in table 14.12. No exact test for this kind of relationship is at present known. By analogy with the F -test, one suggestion is to use the ratio

$$\frac{s_t^2 + s_{tpy}^2}{s_{tp}^2 + s_{ly}^2}$$

where the s^2 values are the mean squares in the analysis of variance. This ratio does not follow the F -distribution, but following an approximation suggested by Satterthwaite (14.7) and others, we might use the F -tables with n_1' and n_2' degrees of freedom, where

$$n_1' = \frac{(s_t^2 + s_{tpy}^2)^2}{\frac{s_t^4}{n_t} + \frac{s_{tpy}^4}{n_{tpy}}}$$

where the n 's are the numbers of degrees of freedom in the corresponding mean squares. The analogous expression is used for n_2' .

After completion of the analysis, the next step is to examine the nature of the interactions and average effects of the treatments, bringing to bear any external knowledge that will throw light on the interpretation. Often, for the practical uses of the results, we would like to decide whether a single recommendation can be made for application in the whole population, or whether, on the other hand, it is necessary to have several recommendations for different parts of the population. These decisions are facilitated by calculating confidence limits for the differences among the means of the most likely candidates. For a discussion of methods for making this calculation, which can give rise to some complications, see reference (14.5).

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CHAPTER 15

RANDOM PERMUTATIONS OF 9 AND 16 NUMBERS

15.1 Use of the Random Permutations

In practically all the experimental designs, the randomization consists in arranging a set of objects, whether treatments, blocks, rows, or columns, in random order. If the number of objects does not exceed 9, a randomization is obtained at once from table 15.6, which contains 1000 random arrangements of the numbers from 1 to 9. For example, to arrange 7 treatments in random order, select a starting place in table 15.6 (without inspection of the numbers in the table). Suppose that the permutation chosen is

1 8 9 2 6 7 5 3 4

Omitting the digits 8 and 9, we obtain

1 2 6 7 5 3 4

for the desired arrangement. Similarly, table 15.7, which contains 1000 permutations of the numbers between 1 and 16, may be used for any number of objects between 2 and 16.

Occasionally the randomization involves dividing the objects into groups, as in the completely randomized and cross-over designs. Thus, with a cross-over design (section 4.4) having 2 treatments and 10 blocks, treatment *A* must appear in the first row in five of the blocks, chosen at random, and in the second row in the remaining five blocks. Select a random permutation of the numbers up to 16 from table 15.7, say

6 16 10 5 12 7 9 4 1 2 15 14 13 11 8 3

The numbers from 11 to 16 are ignored. The first five of the remaining numbers, 6, 10, 5, 7, and 9, are chosen as the blocks in which *A* appears in the first row.

If the number of objects to be randomized exceeds 16, see section 15.3.

15.2 Construction of the Random Permutations

The idea of using random permutations instead of the ordinary random digits was suggested by Professor George W. Snedecor. Before the

writing of this book, 200 random permutations of each of the numbers from 3 to 13 had been made at the Statistical Laboratory at Iowa State College. These were obtained by first constructing a large number of ordinary random digits, which were then transformed into random permutations. The method of transformation was the usual one: it may be illustrated for the numbers from 1 to 11. To obtain a random permutation of the numbers 1 to 11, a series of pairs of random digits is taken, say

65, 04, 29, 82, 37, 52, 53, etc.

When we divide each number by 11 the remainders are

10, 4, 7, 5, 4 (omitted), 8, 9, etc.

This process is continued until 10 of the 11 numbers have appeared, repetitions being omitted.

Rather than present 200 permutations of each of a series of numbers, it was decided to concentrate on two numbers. This decision was made partly because any permutation of 9 numbers, say, can be used to give permutations of all numbers less than 9. Thus 1000 permutations of 9 numbers are more useful than 200 of each of the numbers 9, 8, 7, 6, and 5. Also, certain numbers, such as 11 and 13, are less often used in experiments than others. The numbers 9 and 16 were selected for presentation because of their occurrence in factorial experiments.

The 1000 permutations of 9 were obtained from the permutations already available for 9, 10, 11, 12, and 13, by omitting the numbers above 9. Of the permutations of 16, 800 were obtained from random digits. With as many as 16 numbers to permute, the method of division as illustrated above is rather slow. A more rapid alternative, suggested by Mr. Paul Peach, was used. In this method 16 pairs of random digits are taken, for example,

75	10	17	28	91	85	74	56	71	06	26	01	06	21	07	60
14	5	6	9	16	15	13	10	12	3	8	1	2	7	4	11

The permutation is produced by ranking the pairs in order of size, as shown below each pair. In the event of a tie, as happens in this example with the two 06's, the two ranks in question may be allotted from an additional random digit. Ties occur frequently (the probability that all 16 pairs are distinct is about .35) but cause little delay. Although the method has the additional merit of using fewer random digits than the method of division, with 32 random digits per permutation a large supply of random digits is required. The remaining 200 permutations were obtained by drawing numbered marbles from an urn.

the numbers 1, 2, ..., 9 have an equal probability of appearing in the 1st, 2nd, ..., 9th positions. This test is made by counting the number of times that each number has occurred in each position. The results are shown in two-way presentations in tables 15.1 and 15.2.

For the permutations of 9, the expectation in each cell is $1000/9$, or 111.1. Each row and each column in table 15.1 adds to 1000. Taking the values in a single column, we may test by χ^2 whether the 9 numbers are equally represented in the corresponding position. If the x 's are the observed entries in the column,

$$\chi^2 = \sum \frac{(x - m)^2}{m} = \frac{9}{1000} \sum \left(x - \frac{1000}{9} \right)^2$$

For computational purposes, this simplifies to

$$\chi^2 = \frac{9}{1000} \sum (x^2) - 1000$$

Thus, for the first position,

$$\chi^2 = \frac{9}{1000}(130^2 + 120^2 + \dots + 112^2) - 1000 = 10.81$$

with 8 d.f. Similarly, from the rows we may test whether any given number has appeared equally often, apart from sampling fluctuations, in each position. Although the two sets of tests are not independent, both are of interest. The values of χ^2 are shown in table 15.3.

TABLE 15.3 VALUES OF χ^2 FROM 1000 PERMUTATIONS OF 9

Position		Number	
test	χ^2	test	χ^2
1	10.81	1	7.46
2	5.19	2	9.17
3	5.08	3	6.70
4	17.16	4	7.75
5	6.06	5	18.90
6	15.52	6	8.97
7	8.82	7	8.25
8	3.88	8	3.45
9	3.44	9	5.32
Totals	75.96		75.97

In the *position* tests, the individual probability values range from .90 ($\chi^2 = 3.44$) to .028 ($\chi^2 = 17.16$). With 8 d.f., the probability that the largest of 9 independent values of χ^2 should exceed 17.16 works out at .23. Actually, owing to the fact that the cell numbers add to 1000

in any row or column, the 9 values for the position χ^2 have a small positive correlation which reduces the probability value for the largest χ^2 to about .20. The individual probability values for the *number* tests all lie between .90 ($\chi^2 = 3.45$) and .015 ($\chi^2 = 18.90$). The probability that the largest χ^2 should exceed 18.90 is about 0.11. Thus neither series of tests indicates any marked departure from randomness.

The total, 75.97, which is $\sum (x - m)^2/m$ taken over all cells in table 15.1, supplies a composite test of the complete table. On account of the positive correlation referred to in the previous paragraph, the value 75.97 does not itself follow a χ^2 distribution. In fact, it may be shown to be distributed approximately as $(9/8)\chi^2$, where χ^2 has 64 d.f. Consequently we may test $\frac{(8)(75.97)}{9}$, or 67.52, as a χ^2 value with 64 d.f.

As would be expected from the results of the previous tests, the probability value, .36, provides no evidence for the rejection of the null hypothesis.

The corresponding χ^2 values for permutations of 16 are given in table 15.4.

TABLE 15.4 VALUES OF χ^2 FROM 1000 PERMUTATIONS OF 16

Position test	χ^2	Number test	χ^2
1	14.40	1	20.25
2	17.70	2	12.96
3	11.87	3	16.58
4	15.14	4	4.80
5	17.70	5	14.94
6	7.42	6	9.57
7	12.29	7	13.98
8	7.42	8	13.63
9	24.54	9	15.55
10	9.15	10	12.45
11	10.62	11	14.05
12	16.51	12	18.56
13	9.92	13	14.50
14	18.85	14	13.06
15	6.98	15	14.78
16	19.36	16	10.21
Totals	219.87		219.87

The only figure in the group which has an unusual probability is the low value 4.80 for number 4. This has an individual probability of .9932 (15 d.f.). When account is taken of the fact that this is the smallest of

the 16 values for the number tests, the probability is found to be .10. The total, 219.87, is distributed as $(16/15) \chi^2$ with 225 d.f. and is found to be slightly but not abnormally below its expectation.

A further test of randomness was made by counting the number of *inversions* for each permutation. The inversions are counted as follows. For each number, record how many numbers to the left are greater than the number. Thus for the permutation

3 4 8 1 9 7 2 6 5

we record

0 0 0 3 0 2 5 3 4

The total $(3 + 2 + 5 + 3 + 4 = 17)$ gives the number of inversions. The extreme values for the number of inversions are zero (for the permutation 1 2 3 4 5 6 7 8 9) and 36 (for the permutation 9 8 7 6 5 4 3 2 1).

TABLE 15.5 OBSERVED AND THEORETICAL FREQUENCIES OF NUMBERS OF INVERSIONS FOR PERMUTATIONS OF 9

Number of inversions	Observed frequency	Theoretical frequency	Goodness of fit, χ^2
0-6	8	6.3	0.46
7	4	6.0	0.67
8	11	9.9	0.12
9	10	15.3	1.84
10	13	22.1	3.75
11	36	30.4	1.03
12	38	39.7	0.07
13	50	49.5	0.01
14	71	59.1	2.40
15	65	67.8	0.12
16	80	74.6	0.39
17	71	79.0	0.81
18	90	80.6	1.10
19	75	79.0	0.20
20	80	74.6	0.39
21	57	67.8	1.72
22	58	59.1	0.02
23	45	49.5	0.41
24	40	39.7	0.00
25	27	30.4	0.38
26	27	22.1	1.09
27	18	15.3	0.48
28	7	9.9	0.85
29	11	6.0	4.17
30-36	8	6.3	0.46
Totals	1000	1000.0	22.94

In intermediate cases the number of inversions serves as a measure of the extent to which the order is inverted from the natural order towards the reverse order. The frequency distribution of the number of inversions has been studied by Rosander (15.5) and Kendall (15.6), who derived from the number a measure of the rank correlation between two series of figures. For random permutations of 1 to 10 numbers, Kendall also tabulated the exact distribution of the number of inversions. His distribution was extended to 16 figures for the test given here.

For permutations of 9 the observed and theoretical distributions are shown in table 15.5.

The total χ^2 , 22.94, has 24 d.f. and agrees closely with expectation. The corresponding value for permutations of 16 is $\chi^2 = 47.52$ with 54 d.f. and a probability value of .72.

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15.5 Tables of Random Permutations

TABLE 15.6 PERMUTATIONS OF 9

55671	43373	87463	97494	92288	27935	83194
41282	71129	95782	89366	17724	48573	37456
93329	88845	24616	36778	74471	73286	61222
79743	55292	16535	78519	51913	65149	29878
16965	69436	43929	51823	83332	89612	45769
64436	24681	79341	62642	29859	92428	96981
87817	12568	31298	44187	65167	54351	14317
32194	36757	68877	25951	38546	36794	52545
28558	97914	52154	13235	46695	11867	78633
74615	92229	28173	24219	24831	26548	84942
93832	11198	94954	88886	77546	53276	93821
16347	65845	61719	52563	85755	69981	36797
68284	48786	57545	96758	59977	85335	69469
41478	23934	42236	47425	63369	17854	45214
29193	79662	16461	79974	18418	92793	18355
55551	37477	85892	15132	96284	38119	57133
82929	86553	79688	31697	41693	44662	72688
37766	54311	33327	63341	32122	71427	21576
97755	99938	98617	58612	19833	31773	76655
38172	62716	41342	36243	26128	88627	89747
43427	73172	15486	62161	78517	59136	31231
59283	37589	29171	23834	35999	72341	57178
16511	56441	73723	47388	93256	66959	98912
62836	84625	52268	91756	47464	17464	12886
24964	18354	36594	85979	81681	45595	24594
85699	25267	87839	19425	64745	23282	63323
71348	41893	64955	74597	52372	94818	45469
74987	97171	92387	78535	51649	78618	29734
56112	64614	59128	24687	73761	51741	93477
49356	11848	35493	36123	26877	45385	85951
33228	52322	73869	41861	19236	39577	12812
21494	46283	27651	57312	98413	63129	61588
97545	39799	14234	69744	32522	84263	56363
62639	88555	86772	93458	87994	92494	48129
85871	23937	41515	85976	45358	16852	34645
18763	75466	68946	12299	64185	27936	77296
84686	21997	22189	51924	52628	16883	81941
99458	44878	87597	36477	38536	44677	66878
66311	68319	75755	65185	24382	51436	49786
73772	73622	38946	47269	79741	38265	35314
28934	15551	54364	78753	95865	82792	53435
37269	86463	41821	19648	47213	63551	22699
51845	99184	19432	82896	63499	27124	98262
45527	32736	93218	93512	16977	95918	77157
12193	57245	66673	24331	81154	79349	14523

TABLE 15.6 PERMUTATIONS OF 9 (Continued)

8 6 2 2 4	5 4 5 8 3	9 6 5 2 2	4 9 8 2 9	8 6 6 5 6	3 5 6 6 5	1 1 3 9 1
2 2 3 3 9	8 1 8 3 7	6 5 1 3 8	3 6 9 1 6	3 7 4 3 5	9 3 2 5 2	5 8 1 3 5
7 5 6 9 5	4 2 6 6 1	7 4 4 9 6	1 3 3 7 5	9 4 8 8 8	5 4 1 8 7	2 2 8 8 2
5 4 4 5 3	7 5 2 2 4	5 9 9 8 4	5 8 5 9 4	6 1 2 7 7	1 1 5 4 9	4 4 5 7 9
6 8 1 8 2	9 6 1 4 5	4 7 8 6 9	7 4 2 8 2	2 5 5 4 4	4 8 8 1 3	9 7 6 6 8
1 9 9 1 1	3 9 4 1 9	3 3 3 7 5	6 5 1 4 1	4 8 9 1 1	2 9 9 3 8	8 6 2 5 7
9 1 8 4 8	1 7 9 7 6	2 1 7 4 3	8 2 7 3 7	7 9 1 6 9	6 6 4 2 6	6 5 7 2 4
3 3 5 6 6	2 8 7 5 2	8 2 6 1 7	2 7 4 6 3	5 3 7 9 2	8 7 7 9 1	3 3 4 1 3
4 7 7 7 7	6 3 3 9 8	1 8 2 5 1	9 1 6 5 8	1 2 3 2 3	7 2 3 7 4	7 9 9 4 6
6 5 3 4 8	9 5 3 8 2	1 6 4 4 6	8 4 9 9 1	4 3 3 7 9	9 3 4 5 9	3 2 5 6 1
2 3 7 5 9	6 1 7 5 4	8 4 5 9 1	3 7 1 8 5	7 1 9 1 3	6 9 8 1 5	6 4 4 9 2
7 4 9 8 2	4 7 2 1 9	6 7 2 8 7	5 5 6 6 2	8 8 4 3 7	2 5 7 3 1	5 5 3 2 5
5 2 2 9 6	7 2 5 9 5	3 2 6 1 8	6 2 4 3 9	6 5 6 2 1	3 4 5 6 3	9 3 2 4 4
4 7 5 1 7	2 9 8 6 8	5 9 9 2 5	9 3 2 2 8	2 9 2 4 8	4 2 6 4 2	7 9 7 5 6
8 9 4 3 3	3 6 1 2 7	2 8 8 6 9	4 8 7 7 4	1 2 5 9 4	7 6 1 2 7	2 8 6 7 3
1 1 6 2 1	8 4 6 4 1	9 1 1 3 2	2 1 8 5 7	3 7 8 6 2	5 8 2 9 6	4 7 1 3 7
3 8 1 6 4	1 3 9 3 3	7 5 3 5 4	1 9 3 1 6	5 6 1 5 5	8 7 3 8 4	1 1 9 1 8
9 6 8 7 5	5 8 4 7 6	4 3 7 7 3	7 6 5 4 3	9 4 7 8 6	1 1 9 7 8	8 6 8 8 9
7 4 2 4 2	1 2 2 6 2	8 6 5 2 2	6 1 8 4 7	1 2 1 8 3	9 7 7 4 5	4 6 1 4 7
8 5 9 8 7	8 3 9 2 3	1 1 6 5 9	7 7 2 9 3	5 4 4 7 8	3 4 1 1 7	6 5 8 1 6
9 6 4 5 9	3 1 3 7 5	7 4 2 8 8	9 9 4 5 8	2 9 6 1 7	4 6 3 3 6	9 8 3 7 8
1 7 7 6 3	6 6 7 4 8	9 7 1 6 7	1 5 7 8 5	6 5 2 4 6	6 2 5 9 8	1 4 5 9 2
2 2 8 7 5	5 9 1 5 9	5 5 9 4 6	2 3 3 6 4	7 6 5 3 4	2 5 6 7 4	7 1 9 5 4
5 9 1 2 1	4 8 5 1 6	6 9 3 9 4	3 4 9 2 9	9 1 9 2 1	7 9 2 6 1	3 3 2 2 9
6 3 6 1 6	7 7 8 9 4	3 8 4 1 3	5 6 5 1 1	8 8 8 9 2	5 3 4 5 2	5 7 7 8 5
4 1 5 9 8	2 5 4 3 7	4 3 8 3 5	8 2 6 3 6	4 7 7 6 9	8 8 9 2 3	2 2 4 6 1
3 8 3 3 4	9 4 6 8 1	2 2 7 7 1	4 8 1 7 2	3 3 3 5 5	1 1 8 8 9	8 9 6 3 3
9 7 5 9 7	3 2 3 5 8	1 1 7 6 1	2 2 6 6 3	7 3 8 2 7	4 1 7 6 7	8 9 4 3 7
2 5 7 7 5	2 6 6 7 2	2 7 8 1 3	4 8 8 5 9	8 1 9 8 9	7 9 3 8 8	9 2 1 9 2
7 2 3 3 9	5 1 7 4 4	5 3 2 7 6	7 3 2 3 2	3 8 6 6 5	8 3 8 2 6	5 6 2 8 1
4 4 8 5 2	4 9 5 1 9	8 2 4 5 5	9 9 9 4 5	1 4 3 4 6	9 5 6 9 3	7 8 8 4 9
5 3 9 8 8	1 5 2 9 6	6 4 5 9 4	5 5 5 7 7	9 6 2 9 1	6 8 4 4 2	3 7 7 6 8
6 6 1 4 6	9 3 1 2 7	4 8 9 4 2	6 7 1 2 8	4 7 7 7 3	5 4 9 3 9	1 1 6 1 5
3 9 4 6 4	7 4 9 3 5	9 9 6 2 8	3 1 7 8 4	6 2 5 5 4	1 6 1 7 4	4 4 9 2 6
8 8 2 2 1	6 7 8 6 3	7 6 1 8 7	8 6 4 9 1	2 9 4 1 8	3 7 5 5 1	2 3 3 7 4
1 1 6 1 3	8 8 4 8 1	3 5 3 3 9	1 4 3 1 6	5 5 1 3 2	2 2 2 1 5	6 5 5 5 3
2 5 6 6 5	4 2 8 3 2	4 8 6 2 6	4 2 4 1 3	9 8 1 9 7	4 7 4 9 6	1 3 4 1 4
8 1 1 5 7	6 5 7 1 8	7 7 2 7 8	3 1 1 6 8	3 7 2 6 5	8 5 6 7 2	4 1 9 5 9
4 9 2 3 9	3 6 6 6 6	3 4 3 4 7	5 8 2 3 5	8 1 6 7 1	3 2 8 2 4	3 2 8 7 3
7 8 4 8 1	9 4 4 9 3	5 6 8 3 5	1 9 3 9 6	6 2 8 3 2	5 3 1 6 9	5 4 7 6 5
5 4 5 7 2	2 3 9 4 5	9 9 4 8 2	9 3 5 4 9	1 5 3 4 4	6 4 9 4 8	2 5 3 4 1
3 2 3 9 6	1 8 1 7 7	6 5 1 6 9	7 5 6 2 4	7 6 7 8 8	1 1 5 8 3	6 7 1 3 6
1 6 9 2 3	5 7 3 2 9	2 3 7 1 1	6 7 7 7 1	5 3 5 5 6	9 8 3 3 1	7 6 6 9 8
6 3 7 1 8	8 9 2 8 4	1 1 9 9 3	2 4 9 5 7	2 4 9 2 9	2 6 2 1 7	9 8 5 2 7
9 7 8 4 4	7 1 5 5 1	8 2 5 5 4	8 6 8 8 2	4 9 4 1 3	7 9 7 5 5	8 9 2 8 2

TABLE 15.6 PERMUTATIONS OF 9 (Continued)

73653	32896	31468	84686	63712	58535	18475
55426	91537	22274	66364	78368	94979	72957
48815	49759	16912	29199	46293	87428	94523
37348	55121	88559	51522	37574	61114	57848
89597	14912	64687	33971	59155	13646	33639
26981	28363	47821	48413	14437	79287	49261
64232	77685	53393	72235	82946	32362	65194
91179	63448	95745	15858	21829	26751	26382
12764	86274	79136	97747	95681	45893	81716
47712	44429	87455	71191	22417	88271	47698
94224	21664	52337	43589	97793	26848	89181
31836	53988	43178	84268	75838	45792	74752
18347	87231	95823	25374	39956	53134	53523
53998	96596	28244	17842	58684	62566	22914
62571	68847	31696	96927	86549	94955	95269
26189	35353	74712	38636	44161	79319	68475
79655	72112	69561	59413	61375	37687	16336
85463	19775	16989	62755	13222	11423	31847
74785	35957	67413	22284	42225	44417	11517
17546	54176	88385	51956	37751	58694	85133
26119	88833	26596	95845	64683	19721	43264
63627	77415	93131	33171	95137	95953	69856
35498	26669	42957	67737	56392	63272	58791
48962	99782	39779	16599	21569	81168	37625
91251	41241	74644	48413	13446	22536	22488
52834	62328	51222	84662	78874	27349	94379
89373	13594	15868	79328	89918	76885	76942
11289	72161	91142	71644	61991	79225	61949
67153	49644	84687	92426	95222	63359	55261
73975	56786	35875	59962	49676	46161	44852
39512	31259	66731	86553	74559	57787	37475
54491	28532	23224	14185	36733	34816	29198
85836	15977	52519	38718	83818	15444	93336
92768	83828	19398	45297	58164	21593	82723
46344	64313	47453	63339	27345	88938	76587
28627	97495	78966	27871	12487	92672	18614
37415	52698	87644	17966	71833	33716	89525
78962	21365	96222	71459	93211	27565	62949
24649	45486	68316	95111	29765	19348	55117
46536	19533	32767	24348	16149	76893	93281
12771	34951	75971	56775	54627	45979	21872
59897	73117	59185	33882	68954	88124	18498
65323	97772	48438	62537	37478	62657	76666
91184	88829	24599	48223	82396	51432	47333
83258	66244	11853	89694	45582	94281	34754

TABLE 15.6 PERMUTATIONS OF 9 (Continued)

9 8 1 1 9	4 7 6 3 4	6 2 1 2 8	7 4 8 2 4	2 6 3 1 6	6 9 9 6 7	9 9 2 4 2
4 2 2 9 3	6 2 7 8 1	3 9 6 3 7	5 6 9 4 5	9 3 6 6 1	3 5 1 5 3	2 6 8 3 7
7 1 9 2 6	1 9 5 6 3	5 8 8 7 3	4 1 6 1 1	1 2 1 9 4	2 4 2 2 8	1 7 7 9 8
1 7 4 5 5	5 8 8 5 7	1 1 7 6 4	1 9 4 5 2	5 7 9 7 5	4 7 8 1 5	5 2 5 2 3
6 6 8 3 4	2 5 2 4 5	2 7 2 8 5	2 5 2 9 9	7 1 7 8 2	8 8 6 7 9	3 4 1 1 4
2 9 6 6 2	8 3 1 9 6	9 3 5 1 6	3 2 7 7 7	6 4 8 4 3	9 2 5 8 1	7 3 3 7 5
3 5 3 4 1	7 4 9 1 8	4 4 9 4 9	9 3 1 8 8	8 5 4 2 8	7 1 7 4 2	6 8 4 8 6
8 4 7 8 7	9 6 4 7 9	7 6 4 9 1	6 8 5 6 3	3 8 2 5 9	1 6 3 9 6	8 1 6 5 9
5 3 5 7 8	3 1 3 2 2	8 5 3 5 2	8 7 3 3 6	4 9 5 3 7	5 3 4 3 4	4 5 9 6 1
2 4 8 1 4	9 9 9 5 2	5 6 3 7 8	3 2 3 8 1	2 1 1 4 8	9 7 2 9 7	7 2 8 4 8
5 2 4 9 8	8 7 3 8 3	2 2 1 3 1	6 9 9 1 9	1 5 4 6 6	3 6 8 6 6	9 8 7 9 4
7 1 6 7 5	7 8 5 3 6	7 3 4 9 5	2 7 7 7 8	7 7 6 2 2	5 5 1 5 9	5 1 2 6 3
9 8 5 8 1	3 3 1 6 4	4 9 7 6 9	8 6 2 5 7	8 8 2 5 5	7 2 9 2 8	8 5 4 2 6
3 7 1 3 7	4 5 7 7 5	9 7 9 1 3	5 1 4 3 5	9 3 9 1 3	1 4 3 1 2	1 6 9 7 5
6 3 2 2 6	1 6 6 9 1	3 8 5 8 6	9 3 1 2 2	5 4 8 7 4	2 9 5 8 1	4 4 1 1 2
8 6 3 4 9	6 4 8 4 9	6 1 6 2 2	1 5 6 4 4	6 2 3 3 1	6 3 6 4 5	3 9 6 8 1
1 9 7 5 2	2 2 2 1 7	8 5 8 4 7	4 8 8 9 3	4 6 5 9 7	4 8 7 7 4	6 3 5 5 7
4 5 9 6 3	5 1 4 2 8	1 4 2 5 4	7 4 5 6 6	3 9 7 8 9	8 1 4 3 3	2 7 3 3 9
3 5 9 5 1	4 1 8 8 6	6 5 4 5 5	3 9 8 6 3	8 6 8 9 1	2 6 5 3 1	1 2 7 7 2
6 2 7 3 7	2 8 6 2 2	9 7 7 7 4	9 4 3 9 9	6 5 6 1 5	7 1 3 8 5	3 6 3 9 6
9 3 2 8 9	5 3 4 9 1	2 1 3 9 7	1 6 2 1 2	9 8 5 3 2	1 2 4 6 3	8 3 5 2 7
5 7 6 9 3	7 5 7 4 3	4 9 6 6 1	7 7 7 4 7	5 4 3 5 8	4 4 1 2 4	5 9 2 1 3
7 4 8 6 2	8 7 5 1 5	5 4 2 4 6	4 3 5 8 5	1 9 7 7 9	8 7 8 5 9	6 4 9 5 9
1 9 1 7 4	3 9 2 5 9	3 3 1 1 3	8 2 1 5 1	4 7 9 2 4	5 9 6 4 2	7 5 8 3 4
8 6 5 1 8	6 6 1 6 8	1 2 8 3 8	2 5 9 7 4	3 1 1 4 7	9 3 9 9 8	4 1 1 4 1
4 8 4 2 6	1 4 9 3 7	8 8 5 2 2	6 8 6 2 8	7 3 4 8 3	3 8 2 1 7	9 7 4 6 8
2 1 3 4 5	9 2 3 7 4	7 6 9 8 9	5 1 4 3 6	2 2 2 6 6	6 5 7 7 6	2 8 6 8 5
4 6 6 2 2	2 2 5 6 5	5 7 5 6 4	6 2 7 1 6	4 8 3 4 6	2 2 5 7 5	7 6 3 5 6
3 2 2 3 4	7 7 9 7 9	6 1 2 9 1	1 9 1 4 3	1 9 2 2 2	9 8 3 1 3	5 4 1 2 7
2 5 5 5 3	6 6 1 2 7	8 9 6 5 6	4 7 9 3 1	8 4 5 7 7	4 6 7 8 1	4 3 7 1 8
1 4 8 7 1	3 3 8 1 3	4 8 8 7 9	3 3 2 2 9	3 7 7 6 9	7 9 6 9 8	3 8 5 9 9
9 1 4 8 5	9 8 3 8 4	2 3 4 4 8	7 5 5 8 2	9 5 1 1 8	6 7 8 6 2	8 2 8 6 5
5 7 9 9 7	1 5 6 5 6	3 6 1 8 5	5 6 8 6 4	2 1 8 8 4	5 3 9 4 6	6 1 4 8 4
7 8 3 6 6	8 4 2 9 2	7 4 7 2 2	2 8 6 5 7	7 6 6 9 3	8 4 1 3 7	9 9 6 3 2
6 3 1 4 8	4 9 4 3 8	1 5 9 3 7	8 1 4 9 8	5 2 4 5 5	3 1 2 2 9	1 7 2 7 3
8 9 7 1 9	5 1 7 4 1	9 2 3 1 3	9 4 3 7 5	6 3 9 3 1	1 5 4 5 4	2 5 9 4 1
8 5 3 9 2	1 7 9 9 6	5 8 8 8 5	3 8 2 4 7	8 4 1 3 8	7 1 1 6 5	4 4 7 2 2
7 2 5 7 5	9 9 4 7 7	9 1 1 1 7	9 3 8 5 6	7 7 3 4 7	8 2 8 7 2	2 9 1 4 7
5 1 4 5 7	7 2 3 4 1	7 2 3 9 4	4 7 9 1 9	6 2 5 1 9	3 4 7 3 1	8 2 8 9 8
9 6 7 2 4	4 6 8 1 5	2 3 9 3 1	7 5 7 8 5	9 5 7 9 4	1 5 9 2 3	5 7 2 1 3
4 8 6 8 3	2 8 6 2 4	4 6 5 7 8	5 2 1 6 8	1 1 9 8 3	9 9 4 8 8	6 1 5 8 6
6 4 9 6 8	5 1 1 8 3	6 4 7 6 3	1 9 3 3 2	3 3 6 2 2	2 7 2 9 9	7 3 3 5 5
2 7 8 4 6	6 4 5 6 9	8 5 2 5 6	8 1 4 7 1	4 9 4 6 1	5 8 6 1 7	9 5 6 3 4
1 9 2 1 1	3 5 2 3 2	1 9 4 4 9	2 6 6 2 4	5 8 2 5 6	6 6 3 5 6	1 8 4 6 1
3 3 1 3 9	8 3 7 5 8	3 7 6 2 2	6 4 5 9 3	2 6 8 7 5	4 3 5 4 4	3 6 9 7 9

TABLE 15.6 PERMUTATIONS OF 9 (Continued)

4 2 6 5 9	1 4 9 7 8	7 4 6 4 3	2 1 2 2 4	3 3 6 8 1	4 7 1 6 4	9 9 3 2 3
6 8 1 3 1	9 6 4 4 2	2 1 8 3 9	5 9 6 5 9	2 2 7 1 8	7 9 8 9 5	2 4 2 5 4
3 6 4 7 8	7 5 1 8 4	9 2 2 7 8	7 4 4 7 8	1 9 9 2 4	6 3 7 4 9	6 1 5 6 6
1 4 8 2 4	3 7 5 5 6	3 5 9 8 2	6 3 7 3 7	4 5 5 3 9	5 6 2 5 2	4 6 1 3 2
7 9 9 6 6	6 1 7 1 3	5 7 5 6 1	8 5 3 9 3	5 4 4 9 5	3 8 9 7 8	7 3 7 7 1
9 5 3 1 3	4 3 3 2 7	1 6 4 1 5	4 2 8 8 1	9 7 2 4 2	8 4 5 8 6	5 8 4 8 8
2 1 5 8 5	8 2 6 3 5	4 3 1 5 4	1 6 5 4 5	8 6 3 6 3	9 1 3 3 7	8 2 6 9 5
5 3 7 9 7	2 8 8 9 9	6 9 7 2 7	3 8 1 1 2	6 1 1 5 7	1 2 6 1 1	3 7 8 4 7
8 7 2 4 2	5 9 2 6 1	8 8 3 9 6	9 7 9 6 6	7 8 8 7 6	2 5 4 2 3	1 5 9 1 9

1 6 3 8 3	7 2 6 7 9	7 8 1 6 5	1 1 4 4 8	1 2 7 8 1	8 9 7 6 9	7 5 8 1 7
2 2 8 9 8	2 5 5 2 6	3 4 8 5 1	4 8 7 2 1	6 5 1 2 2	9 5 1 4 2	3 9 4 3 8
8 7 7 5 1	9 7 9 5 3	4 3 7 7 9	5 5 2 9 6	2 7 9 5 6	5 3 8 9 4	6 2 6 8 3
3 5 6 1 4	3 6 8 9 1	1 5 4 1 4	8 7 1 7 5	8 8 8 6 7	7 2 9 7 8	8 4 7 7 4
9 3 2 7 5	4 1 7 3 7	9 1 9 3 7	2 4 9 3 4	7 9 4 7 8	3 6 2 5 6	1 7 9 9 1
4 4 9 4 6	1 8 2 1 2	2 2 6 8 2	7 9 3 6 3	4 6 6 1 5	2 4 4 2 5	2 6 2 5 6
7 8 1 6 7	5 4 3 6 8	6 9 3 2 6	3 6 5 8 7	3 4 3 4 9	6 8 6 1 7	9 1 3 6 5
6 1 4 3 9	8 9 4 4 5	5 6 2 9 3	9 2 6 1 2	9 1 5 9 4	4 1 5 8 1	5 8 1 2 9
5 9 5 2 2	6 3 1 8 4	8 7 5 4 8	6 3 8 5 9	5 3 2 3 3	1 7 3 3 3	4 3 5 4 2

4 9 6 6 1	1 1 8 3 1	3 7 5 4 9	9 7 4 9 9	9 4 8 8 3	3 2 5 1 3	9 5 6 8 8
5 3 1 9 6	6 8 2 5 9	6 5 4 9 2	2 8 5 6 3	2 1 9 4 2	8 6 4 2 6	8 1 7 9 6
8 6 8 5 7	7 5 1 1 3	7 3 9 2 7	6 9 7 3 6	8 6 5 1 1	9 5 9 9 8	7 3 8 1 1
3 7 4 4 2	2 2 9 4 5	9 1 3 3 8	1 2 1 1 7	3 9 6 2 9	4 8 2 5 4	1 2 3 7 7
2 8 3 3 9	4 9 6 7 2	8 6 7 8 3	8 1 9 2 8	1 7 3 5 6	5 3 3 3 1	2 9 9 6 9
9 4 7 1 4	3 3 3 8 6	2 9 1 1 4	3 6 3 7 1	4 2 1 3 4	6 9 8 7 5	5 4 2 5 5
1 5 9 2 8	9 6 5 6 8	1 4 6 6 1	5 5 2 5 2	7 5 4 6 7	1 1 1 8 9	4 7 4 3 2
7 1 5 8 3	5 4 7 9 4	5 8 8 7 5	7 4 8 8 5	5 3 7 9 5	2 7 7 6 7	3 8 5 4 4
6 2 2 7 5	8 7 4 2 7	4 2 2 5 6	4 3 6 4 4	6 8 2 7 8	7 4 6 4 2	6 6 1 2 3

3 1 7 2 7	5 4 3 6 3	9 8 6 4 4	8 6 6 9 6	5 8 1 2 6	5 4 1 1 1	1 2 1 7 3
2 2 5 8 8	9 6 5 5 5	3 1 4 8 8	3 9 3 1 7	7 3 7 5 7	6 7 4 4 9	3 7 3 3 4
1 5 8 6 9	2 2 1 2 4	4 9 9 9 1	1 3 4 6 8	8 4 6 7 4	2 8 3 9 2	8 9 5 9 2
6 3 2 7 6	8 5 8 8 1	7 5 7 2 2	4 5 2 5 1	1 2 5 6 5	7 2 9 7 6	4 4 2 4 7
9 8 4 1 4	7 3 7 3 8	6 4 5 3 9	5 7 7 2 9	3 6 2 9 9	4 6 5 2 7	7 6 4 8 1
5 7 6 3 3	4 1 2 7 9	5 2 2 7 7	9 4 1 4 4	2 1 3 3 1	1 9 2 6 3	2 3 8 5 6
4 6 1 5 5	1 7 4 4 6	1 3 1 1 5	6 8 9 8 3	6 7 4 4 8	3 3 8 5 5	9 8 6 6 8
8 4 9 9 1	3 9 6 9 2	8 6 8 5 3	2 1 5 7 5	4 5 9 1 2	8 5 7 3 8	5 1 7 1 9
7 9 3 4 2	6 8 9 1 7	2 7 3 6 6	7 2 8 3 2	9 9 8 8 3	9 1 6 8 4	6 5 9 2 5

9 2 4 9 5	4 8 4 4 8	1 9 4 8 5	2 7 9 6 5	9 8 7 3 4	3 8 2 1 3	3 5 3 2 6
1 1 8 1 3	8 6 5 9 9	2 7 6 7 7	6 8 6 9 8	2 2 2 2 9	1 4 8 6 2	2 8 9 8 4
3 9 5 5 7	2 4 9 3 3	8 1 9 2 3	7 6 5 7 7	6 7 8 6 7	2 5 9 5 7	1 4 1 1 8
8 6 6 7 2	6 9 1 5 6	9 6 5 3 1	1 1 7 5 1	8 3 4 5 8	9 3 4 2 8	5 1 6 7 2
7 5 3 3 9	7 3 6 8 7	6 8 2 5 4	3 4 1 4 6	5 9 5 9 2	6 2 5 7 5	6 9 7 3 7
2 4 9 2 1	9 7 2 2 4	4 2 7 4 8	8 3 4 3 2	4 6 9 7 1	7 7 6 9 6	7 2 2 6 1
4 3 7 8 4	5 1 8 1 2	7 3 3 9 9	9 9 2 1 9	3 1 1 1 3	8 9 7 3 4	4 3 5 4 9
6 8 1 6 6	3 5 7 7 1	5 5 1 1 6	5 2 3 8 3	1 5 6 8 6	4 6 3 8 9	8 6 4 9 5
5 7 2 4 8	1 2 3 6 5	3 4 8 6 2	4 5 8 2 4	7 4 3 4 5	5 1 1 4 1	9 7 8 5 3

TABLE 15.6 PERMUTATIONS OF 9 (Continued)

2 3 1 7 1	4 2 4 3 1	9 2 1 1 1	4 6 9 3 5	3 6 2 5 5	1 1 4 1 2	7 1 4 5 3
5 9 6 3 3	3 4 6 8 6	4 1 8 6 6	8 1 2 7 9	1 5 1 2 8	6 6 3 4 5	5 6 1 4 6
8 4 5 5 8	6 1 9 1 5	5 6 9 4 7	6 3 5 4 7	5 1 7 7 2	3 8 1 5 4	3 2 2 8 7
1 6 8 1 7	2 9 2 5 3	8 9 4 7 9	7 4 7 2 2	9 4 9 9 3	9 5 9 2 1	9 8 6 7 4
4 1 2 4 9	5 5 3 6 9	7 8 6 3 8	1 8 4 9 6	6 2 8 1 7	4 7 8 3 7	8 3 3 9 8
6 2 7 8 6	8 3 7 4 7	2 5 7 5 3	3 2 6 8 8	2 8 5 3 9	2 2 2 9 6	1 7 8 2 2
3 5 4 9 2	1 7 8 7 4	3 4 2 8 5	5 9 8 6 4	8 3 3 8 4	5 4 6 8 3	4 9 5 6 9
9 8 3 6 5	7 6 5 2 2	1 7 5 9 2	2 7 1 5 3	7 9 4 6 6	8 3 7 7 8	6 4 9 3 1
7 7 9 2 4	9 8 1 9 8	6 3 3 2 4	9 5 3 1 1	4 7 6 4 1	7 9 5 6 9	2 5 7 1 5
9 9 6 8 1	9 5 1 6 2	8 9 5 8 9	7 7 4 9 9	4 9 5 4 6	5 9 5 5 2	6 8 9 7 1
1 7 9 5 6	1 8 7 5 1	1 1 7 7 8	2 4 7 4 7	9 5 1 2 2	1 6 1 9 8	8 6 6 1 8
7 3 2 7 9	4 1 2 4 3	3 6 1 3 3	3 3 8 6 1	5 6 3 5 5	4 4 4 2 9	5 3 1 3 7
8 5 7 4 4	5 9 4 1 4	7 2 3 4 6	5 9 2 2 6	3 8 8 1 1	7 8 7 1 4	4 1 8 9 9
4 8 8 6 8	3 2 3 9 7	6 7 2 5 1	6 6 3 8 5	2 1 4 3 3	3 7 6 7 7	1 9 7 6 2
6 1 4 9 7	8 3 9 8 6	5 8 6 1 2	8 2 5 1 2	1 4 2 9 8	9 2 2 8 3	2 7 2 4 5
2 6 1 3 2	6 7 5 3 5	2 5 4 6 7	1 1 6 5 3	6 7 7 8 9	8 3 3 6 5	3 5 3 8 3
3 2 5 1 3	7 6 6 2 8	9 3 8 2 5	9 8 1 3 8	8 3 6 7 4	6 1 9 3 6	7 4 4 2 6
5 4 3 2 5	2 4 8 7 9	4 4 9 9 4	4 5 9 7 4	7 2 9 6 7	2 5 8 4 1	9 2 5 5 4
9 1 7 8 7	9 8 6 1 4	2 9 1 7 7	4 8 4 8 6	8 2 3 8 2	8 2 3 2 6	7 3 8 7 5
4 8 9 6 3	3 7 9 2 1	9 5 9 4 4	9 4 3 4 3	7 3 5 7 6	5 9 4 6 8	5 8 4 5 6
7 4 2 2 2	6 3 3 6 9	3 1 5 2 1	6 9 9 9 7	5 6 2 6 9	2 6 6 8 1	8 6 5 1 8
5 9 1 3 4	1 9 7 5 3	6 2 8 6 5	1 7 8 7 9	1 4 4 9 8	9 1 7 3 9	1 9 6 6 2
3 3 3 7 5	8 4 2 7 6	1 6 3 5 6	3 6 1 6 1	9 1 1 5 5	6 7 9 4 4	3 1 3 8 7
2 2 5 1 1	7 5 8 8 8	5 7 2 8 3	8 3 6 3 8	4 7 6 4 3	1 3 5 1 3	9 2 1 4 9
1 6 8 5 6	2 2 5 3 5	8 3 4 9 8	2 5 2 5 4	2 5 7 3 7	4 5 1 5 2	6 4 2 2 1
8 5 6 4 9	4 6 4 4 7	7 8 6 3 2	5 1 5 1 5	6 8 9 1 4	3 4 2 9 5	4 7 7 9 3
6 7 4 9 8	5 1 1 9 2	4 4 7 1 9	7 2 7 2 2	3 9 8 2 1	7 8 8 7 7	2 5 9 3 4
6 3 8 7 9	5 1 9 8 5	5 1 8 5 4	1 1 2 2 4			
5 4 6 4 8	7 4 2 4 7	9 3 6 8 9	5 2 1 6 8			
9 8 5 3 7	6 7 8 5 6	6 9 4 1 1	7 5 5 3 3			
1 2 1 9 4	8 6 1 1 8	7 5 3 7 8	2 9 9 5 9			
3 7 2 6 1	1 8 6 6 3	4 8 9 4 5	8 4 8 4 5			
4 9 3 1 6	3 5 4 2 2	8 7 7 6 2	9 7 7 8 2			
8 5 4 5 5	9 9 7 9 9	3 4 5 2 7	3 8 3 1 7			
7 6 7 2 3	2 3 3 7 1	2 6 1 3 3	4 3 6 7 1			
2 1 9 8 2	4 2 5 3 4	1 2 2 9 6	6 6 4 9 6			

TABLE 15.7 PERMUTATIONS OF 16

7 12 15 15 1	2 7 16 10 2	14 15 7 13 13	10 6 1 8 10
13 3 8 16 7	10 11 10 13 5	11 7 13 16 7	7 5 13 2 14
3 1 4 5 14	13 3 14 9 13	13 2 9 15 6	2 8 4 5 8
11 8 16 14 15	6 2 6 2 16	8 5 12 3 9	13 4 3 10 4
14 9 1 6 3	9 14 13 8 6	5 8 14 7 3	15 13 11 4 7
2 16 10 13 5	5 13 2 11 7	3 12 5 14 12	16 2 2 9 15
4 6 13 7 2	15 1 9 1 4	7 10 6 9 11	9 7 6 16 11
6 14 6 10 4	14 4 15 3 3	4 16 2 6 5	1 12 10 6 9
10 15 2 1 13	12 16 3 4 8	10 1 15 5 14	12 14 12 3 2
12 10 7 12 9	11 9 8 12 14	15 4 11 8 16	8 9 14 14 1
15 7 5 2 10	7 8 12 6 15	6 13 16 12 15	4 11 8 12 6
16 2 11 8 8	8 15 5 16 1	1 9 8 1 8	14 16 5 13 5
9 13 14 3 6	4 10 11 5 12	9 3 10 4 4	3 10 9 1 3
8 11 9 4 11	3 12 7 7 10	12 14 3 10 1	6 15 16 15 12
1 5 12 11 16	16 5 4 14 9	16 11 1 2 10	5 1 15 7 13
5 4 3 9 12	1 6 1 15 11	2 6 4 11 2	11 3 7 11 16
11 8 16 5 5	13 1 13 2 16	14 12 9 8 7	5 13 3 13 3
2 2 8 8 14	16 4 3 8 11	10 14 15 1 2	11 4 5 15 9
6 13 2 13 6	5 9 15 11 10	12 6 16 15 16	9 10 12 16 15
14 12 4 16 16	11 14 10 5 12	3 3 12 14 15	13 6 4 1 16
8 6 3 9 4	10 6 4 16 2	2 9 8 16 4	6 5 15 7 8
9 15 12 10 3	2 12 6 1 15	4 13 7 7 9	12 14 8 8 11
3 10 11 12 13	12 5 11 7 8	9 5 14 11 10	1 3 13 3 5
16 1 13 14 8	14 15 5 3 7	11 15 6 12 5	7 11 1 14 4
1 14 14 2 9	15 16 14 6 14	7 8 3 13 11	8 7 7 12 7
4 4 6 4 12	3 11 8 15 9	8 1 13 6 3	3 15 9 9 12
15 5 1 11 10	6 3 7 10 5	5 11 10 10 12	15 16 14 5 2
5 3 5 6 7	7 13 2 14 3	16 4 5 5 13	4 9 16 2 6
12 7 15 15 15	9 8 12 12 13	15 10 1 4 6	16 2 6 11 1
10 11 10 3 2	4 2 1 4 6	6 7 11 9 14	10 8 11 4 13
7 9 7 7 11	1 7 16 13 1	13 2 4 2 1	2 12 2 10 14
13 16 9 1 1	8 10 9 9 4	1 16 2 3 8	14 1 10 6 10
1 6 7 4 8	6 5 2 8 15	4 6 6 1 4	5 7 13 2 10
9 15 11 3 11	15 9 10 1 3	8 2 15 7 9	8 16 1 14 3
10 16 4 5 12	9 16 11 7 1	7 16 11 8 3	3 12 2 3 4
4 14 1 9 5	5 4 13 6 8	15 5 12 5 7	16 5 11 8 1
7 3 13 14 15	2 1 14 16 5	14 9 2 16 1	12 6 14 4 13
16 11 2 1 14	16 6 9 3 4	16 14 3 15 11	11 3 9 12 5
3 10 16 16 13	7 13 1 11 14	9 10 16 2 10	2 10 7 10 16
11 13 9 13 4	13 8 3 5 13	10 12 5 12 5	14 13 16 5 6
15 2 3 12 9	12 2 4 13 10	3 13 14 4 2	1 14 8 6 12
14 1 14 6 10	1 3 12 4 2	2 4 13 3 16	9 9 3 7 14
13 12 5 11 3	11 15 8 2 7	11 7 8 14 6	4 4 4 15 11
12 5 10 7 2	14 7 15 14 16	13 1 9 10 12	10 11 10 9 8
8 9 8 10 6	4 11 7 10 11	6 8 4 9 8	15 8 6 11 9
2 7 6 2 1	8 10 6 15 12	1 11 7 11 13	6 1 15 13 15
6 4 15 8 16	10 14 16 9 6	12 3 10 6 14	7 2 12 16 7
5 8 12 15 7	3 12 5 12 9	5 15 1 13 15	13 15 5 1 2
13 4 10 4 16	13 16 13 5 3	6 14 1 16 8	7 2 3 3 12
5 14 4 6 8	2 15 1 13 14	16 4 15 4 3	12 12 1 4 7
7 12 2 15 14	16 9 12 16 6	10 15 14 9 10	1 14 8 8 16
6 9 7 14 9	3 5 14 7 12	5 13 16 1 7	5 11 2 9 3
14 5 16 7 10	14 10 11 15 11	12 1 12 12 14	16 3 11 11 8
15 11 8 9 7	8 11 8 14 13	7 11 6 3 11	4 4 6 6 9
11 6 6 1 4	12 8 7 1 15	9 3 3 7 13	11 10 4 5 1
4 10 3 16 2	1 3 16 12 5	4 9 13 13 6	8 15 9 1 14
1 8 1 13 1	11 7 9 6 9	1 8 4 11 5	2 16 10 12 4
9 7 14 2 6	15 4 4 11 4	2 16 5 8 1	9 5 12 16 6
12 1 9 10 15	4 14 10 9 8	15 10 7 10 9	10 6 14 10 11
3 3 12 11 5	5 2 15 10 2	14 2 8 2 4	13 8 5 15 5
10 15 11 5 13	9 6 6 3 10	13 12 9 6 2	7 7 15 7 13
8 13 13 3 3	7 12 5 2 7	11 5 10 15 12	3 1 13 13 10
16 16 5 12 11	10 13 2 4 1	8 6 11 14 15	6 9 16 2 2
	6 1 3 8 16	3 7 2 5 16	14 13 7 14 15

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

9	16	15	12	2	11	4	16	11	10	2	5	5	14	11	2	14	13	16	6
11	3	2	6	15	13	10	1	4	13	11	8	16	16	4	3	5	15	5	15
14	14	8	16	11	15	5	14	14	11	1	14	15	15	13	5	7	11	11	16
4	13	1	3	5	7	6	2	16	1	14	9	14	3	3	1	6	16	6	10
6	6	10	7	13	10	16	7	2	12	6	12	6	13	8	9	15	9	1	11
2	10	14	9	12	3	3	10	5	6	5	16	12	10	15	10	11	4	9	8
5	15	11	14	10	4	14	13	6	4	12	4	11	5	10	14	16	5	7	9
16	5	13	10	3	9	12	6	3	7	3	7	3	11	14	7	3	14	4	12
8	12	7	11	7	8	13	15	13	9	4	3	8	1	12	6	9	8	15	14
1	8	3	2	1	5	15	9	9	3	10	11	13	8	5	13	12	3	3	5
13	9	9	1	6	2	11	3	8	8	15	1	7	9	7	8	8	6	2	3
15	1	5	5	9	6	9	4	10	5	8	13	10	7	9	15	2	10	8	4
7	4	12	13	16	1	2	11	12	2	16	15	2	4	2	11	1	7	13	1
10	2	4	15	4	16	1	12	7	15	9	10	9	12	16	4	13	2	10	13
3	7	6	8	8	14	7	5	1	14	13	2	4	2	1	16	4	1	12	7
12	11	16	4	14	12	8	8	15	16	7	6	1	6	6	12	10	12	14	2
12	6	13	4	5	7	2	1	9	2	5	1	15	2	14	13	13	11	2	13
6	11	4	15	12	12	6	15	6	15	6	3	12	5	15	11	16	9	8	1
13	5	1	6	7	6	13	5	7	8	15	6	4	15	1	14	5	14	10	4
11	1	11	7	8	15	8	4	12	13	10	9	3	10	7	2	12	3	9	8
3	7	3	14	15	4	12	11	4	10	8	12	1	4	16	6	2	2	16	7
10	12	15	11	4	13	5	10	3	14	11	2	9	11	2	9	9	12	12	11
15	9	16	16	9	2	16	2	15	6	7	15	8	1	8	12	4	13	6	9
14	15	2	13	3	16	10	14	13	9	10	7	14	9	6	5	6	4	11	12
1	2	12	9	1	8	15	3	8	11	2	5	10	3	3	10	10	7	13	10
5	10	5	3	13	9	9	13	10	1	3	8	7	8	9	4	15	15	7	15
7	14	9	2	11	14	11	6	14	12	9	10	16	12	13	3	7	5	4	14
9	8	10	1	6	3	3	8	5	5	14	16	2	7	12	16	14	10	15	5
2	3	7	5	10	1	1	12	2	7	1	4	6	16	10	8	8	1	5	16
16	13	14	10	2	5	7	16	1	16	13	11	11	6	5	1	11	16	3	3
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8	16	8	12	16	11	4	9	16	4	12	14	5	14	4	7	1	8	1	2
3	14	11	8	9	14	14	2	13	1	8	4	15	16	7	6	15	13	13	13
12	9	6	9	8	10	12	13	14	5	11	10	10	12	9	10	5	16	6	3
11	11	7	1	11	13	11	4	2	7	16	5	8	3	11	12	6	12	5	11
1	16	9	3	1	7	8	15	5	4	3	7	16	8	12	15	7	5	9	4
13	3	1	2	13	5	4	9	7	6	5	15	4	6	4	1	10	6	1	14
7	12	10	10	5	15	5	8	16	2	12	3	5	13	14	13	13	2	3	7
10	15	15	4	14	1	16	16	12	11	9	16	1	2	10	11	8	7	16	8
15	7	4	14	7	4	7	10	6	10	1	1	2	11	3	16	2	4	2	1
9	5	2	7	3	3	13	14	15	15	6	12	9	15	15	9	16	15	15	10
8	6	16	5	15	8	2	12	1	3	10	8	3	14	13	2	1	10	8	13
2	10	5	11	4	9	3	6	11	12	15	9	7	5	2	8	14	1	4	5
5	4	3	15	2	2	15	11	10	14	7	14	14	7	6	3	11	11	10	2
4	1	12	12	16	6	1	3	4	16	13	11	11	4	1	7	12	3	7	9
6	13	14	6	12	16	9	1	8	8	4	13	12	10	5	5	4	9	12	16
16	8	8	13	10	11	10	5	9	13	14	2	6	9	8	14	9	14	11	15
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1	2	14	12	4	4	3	6	12	7	11	11	9	13	13	7	4	10	16	9
9	3	10	13	3	5	5	13	15	9	14	13	14	9	9	4	8	4	15	2
13	6	15	10	11	3	15	12	4	5	5	4	3	6	4	5	12	14	14	3
8	5	5	15	8	9	8	8	2	3	1	12	8	3	11	2	9	16	10	12
11	12	9	14	16	11	4	15	1	4	3	15	5	15	7	11	16	15	7	1
10	4	13	6	1	13	12	9	8	6	7	8	15	7	3	8	13	9	8	10
16	11	11	16	7	15	9	5	7	2	6	10	16	10	6	1	3	6	1	13
5	7	4	3	2	1	14	2	10	13	16	1	6	4	15	6	15	12	11	10
3	15	12	2	14	8	11	16	14	16	9	7	13	8	2	16	2	11	2	15
7	9	7	9	13	6	2	4	13	14	15	6	10	11	8	12	10	3	3	8
2	10	8	8	15	14	6	3	5	1	4	3	7	2	14	15	14	2	6	4
15	8	3	1	6	2	10	7	3	10	10	2	4	1	5	3	7	13	13	14
6	13	2	5	5	7	13	10	16	15	12	16	1	14	16	14	6	1	9	7
12	16	16	4	10	10	7	1	9	8	2	5	12	16	12	9	11	7	4	5
4	1	1	7	9	12	16	11	6	11	8	14	2	5	1	10	1	5	12	6
14	14	6	11	12	16	1	14	11	12	13	9	11	12	10	13	5	8	5	11

TABLE 15.7 PERMUTATIONS OF 16 (*Continued*)

16	2	15	13	13	2	11	14	4	3	12	11	11	11	7	4	13	7	16	16
7	10	8	2	9	5	14	15	5	16	5	9	9	3	1	2	5	6	3	5
9	16	3	8	4	8	7	2	11	1	13	15	2	5	6	8	2	11	6	10
8	11	14	11	14	7	5	7	6	9	16	4	5	16	9	9	16	4	8	15
3	15	11	14	8	10	10	10	12	10	2	2	13	1	11	11	12	13	5	3
12	7	13	15	12	3	12	8	8	11	10	12	10	7	14	15	14	10	12	4
15	13	9	3	3	14	6	16	2	13	8	14	15	14	12	7	1	1	7	13
10	4	1	4	16	6	1	12	16	15	3	16	3	6	16	6	8	3	9	11
6	1	6	5	6	1	15	11	7	8	1	1	8	9	2	16	10	14	14	2
11	8	16	16	5	13	13	6	15	2	15	5	12	8	3	10	11	8	2	14
13	9	10	1	15	15	9	9	3	14	6	6	1	10	13	13	9	5	15	1
14	5	4	6	1	12	3	4	14	6	4	10	4	4	15	3	15	2	1	12
5	12	7	12	10	11	8	3	1	4	7	8	7	15	10	5	3	9	11	7
2	3	2	10	7	4	16	5	10	5	14	13	6	12	5	12	4	16	4	9
4	14	12	7	11	16	4	13	13	12	11	3	14	2	4	14	6	15	10	8
1	6	5	9	2	9	2	1	9	7	9	7	16	13	8	1	7	12	13	6
13	4	3	11	16	8	7	16	3	16	1	10	1	15	16	10	11	4	6	4
4	1	6	16	6	11	11	14	4	8	3	3	5	3	6	7	7	8	5	12
2	7	15	5	7	7	16	3	2	7	10	8	11	11	7	16	12	15	9	9
3	8	12	10	8	6	4	4	11	10	9	2	4	7	1	3	15	10	14	11
6	15	4	6	3	10	2	12	15	6	6	16	9	4	8	12	8	2	8	6
5	6	10	2	13	15	8	7	16	3	2	12	10	9	12	14	9	5	13	5
12	11	1	13	4	5	3	15	5	12	12	1	15	5	15	6	4	12	4	14
9	9	11	8	2	3	12	9	14	13	13	6	8	6	13	11	3	1	10	7
7	13	5	7	1	14	6	13	8	11	14	5	3	16	4	13	10	13	15	3
10	2	14	14	15	16	1	2	1	9	8	15	2	14	9	1	1	7	3	13
14	16	2	1	14	9	13	5	6	4	11	13	16	12	10	15	2	9	2	16
16	3	8	9	5	1	14	6	10	1	16	9	6	10	14	5	6	6	7	15
11	10	16	12	12	12	10	1	12	14	7	14	12	8	3	4	5	16	12	8
15	5	9	15	9	13	9	11	9	5	4	4	13	2	5	8	14	3	1	1
1	12	7	3	10	2	5	10	13	15	5	7	14	1	2	2	16	11	16	2
8	14	13	4	11	4	15	8	7	2	15	11	7	13	11	9	13	14	11	10
1	4	10	3	8	16	4	10	6	10	12	13	16	6	12	10	5	12	1	10
3	10	12	5	7	11	3	3	7	1	6	1	13	2	7	12	16	10	11	13
13	15	5	1	1	12	15	12	14	4	10	14	11	12	4	15	13	16	9	3
10	5	2	9	13	2	16	9	4	2	7	7	4	10	10	6	1	9	5	9
7	6	15	14	15	15	2	7	9	15	2	10	10	8	11	14	4	3	8	15
11	1	11	8	9	7	12	11	11	5	11	3	6	4	15	1	12	4	6	1
12	3	4	2	3	3	1	13	13	12	9	11	1	3	6	2	15	14	7	4
14	9	7	16	14	1	10	16	16	6	8	15	3	16	14	16	14	2	2	7
5	12	13	4	4	14	14	4	5	7	16	5	14	14	8	5	7	15	10	12
4	13	14	13	6	6	8	14	8	9	13	10	7	7	9	3	3	13	12	6
8	14	16	12	12	13	7	2	10	13	4	12	12	5	16	9	11	6	3	16
16	7	3	10	11	9	13	5	12	16	5	9	8	9	3	13	8	8	14	11
6	2	9	15	5	8	9	8	1	11	15	6	9	13	5	11	2	11	13	5
15	8	6	6	16	4	11	1	2	8	3	4	15	15	13	7	6	5	4	14
9	11	1	7	10	5	6	6	15	14	1	2	5	11	1	8	10	1	15	8
2	16	8	11	2	10	5	15	3	3	14	8	2	1	2	4	9	7	16	2
6	12	2	3	9	14	1	16	11	14	5	7	15	11	10	14	10	16	4	16
1	2	15	9	13	15	5	12	12	9	8	6	1	2	5	1	11	13	13	15
9	7	8	13	12	7	11	13	10	5	6	13	6	5	13	9	15	4	9	10
11	1	16	1	1	5	12	4	2	10	16	4	3	16	2	8	9	8	7	7
7	14	11	5	3	4	2	15	5	12	14	9	5	10	15	5	13	9	3	4
8	11	10	10	15	12	8	11	16	13	10	10	10	3	4	12	1	3	2	11
2	4	5	6	16	16	3	2	8	4	7	11	4	7	3	2	7	14	11	1
5	9	7	8	5	6	14	7	14	1	1	2	2	15	7	4	14	15	8	8
16	5	1	11	4	8	15	5	13	8	15	16	8	9	9	11	16	12	12	3
13	10	6	12	6	11	10	6	15	15	2	3	13	1	8	10	5	10	5	12
10	3	14	4	11	3	9	1	3	11	3	14	14	6	12	3	8	1	15	5
12	15	3	14	8	9	6	14	1	6	9	15	11	14	6	7	6	2	6	6
3	16	9	15	2	1	4	3	9	16	13	1	7	13	1	16	3	11	16	13
4	8	4	2	10	10	13	9	6	7	12	8	16	8	11	15	2	6	14	2
15	13	13	16	14	2	16	8	7	3	4	5	9	4	14	13	12	5	1	14
14	6	12	7	7	13	7	10	4	2	11	12	12	12	16	6	4	7	10	9

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

14	15	12	8	7	4	13	14	3	10	13	14	6	3	14	8	10	13	11	14
7	1	8	11	15	11	7	2	12	15	5	8	13	5	6	13	14	2	10	13
13	9	2	15	2	13	6	5	10	3	16	7	5	15	4	7	2	12	4	10
11	3	4	6	11	16	16	4	9	7	11	1	14	9	9	12	3	10	6	2
3	16	3	13	5	1	4	6	14	6	4	3	16	8	15	11	6	1	1	15
5	12	6	9	13	2	12	10	6	1	14	16	11	12	16	3	1	7	12	1
16	5	10	10	12	5	14	15	7	12	15	2	2	10	5	9	7	6	15	7
2	4	16	12	14	3	11	1	13	14	10	6	10	11	13	14	5	15	13	4
6	6	9	3	16	6	2	12	10	13	12	5	4	14	10	15	12	5	3	8
4	7	15	14	6	14	10	9	11	8	8	15	8	13	11	2	9	8	14	12
15	14	5	7	8	10	15	8	4	2	3	11	1	6	2	16	4	4	5	5
9	13	14	4	4	9	5	3	8	9	2	4	3	4	12	5	11	16	8	6
1	2	1	2	1	7	9	11	15	16	1	13	15	2	1	10	15	3	2	9
12	11	13	1	9	8	3	7	2	11	9	9	9	7	3	1	13	11	7	11
8	8	11	16	3	12	1	13	1	5	6	12	12	16	8	6	16	14	16	16
10	10	7	5	10	15	8	16	5	4	7	10	7	1	7	4	8	9	9	3
15	1	3	2	7	5	4	11	1	11	16	13	13	12	15	8	2	7	6	11
13	16	13	7	4	16	7	9	5	1	11	11	5	14	13	5	15	15	4	12
5	14	12	11	14	4	8	7	15	13	6	16	2	1	9	3	9	14	5	6
3	15	9	3	9	11	9	8	12	6	9	14	4	2	7	11	16	5	3	15
6	11	10	9	16	14	2	10	7	14	2	6	16	3	2	4	6	11	8	1
9	2	1	16	13	6	5	5	4	5	5	4	15	16	11	12	4	2	10	7
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14	3	7	4	2	13	16	2	9	7	14	10	14	13	10	2	8	12	13	5
8	12	4	10	5	2	6	1	16	10	1	12	12	9	12	15	10	4	15	2
7	4	16	8	10	9	3	15	2	3	7	2	8	7	1	7	1	1	12	10
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2	13	5	1	3	15	15	3	8	2	10	5	3	15	6	13	7	3	14	3
16	10	11	15	15	7	1	16	13	12	4	9	6	8	8	9	5	9	1	13
12	5	6	14	1	12	10	14	6	15	8	7	10	10	5	16	3	10	16	16
11	6	15	6	11	10	14	4	11	16	12	8	7	11	3	1	14	16	2	8
1	8	14	5	6	8	13	13	14	4	15	3	1	4	14	6	11	6	11	14
9	16	5	1	10	2	8	1	13	11	4	14	6	7	12	15	2	8	8	1
13	7	12	16	7	9	16	12	1	14	1	9	8	10	3	16	9	11	13	13
12	8	16	2	8	8	14	4	3	2	14	7	1	16	4	3	5	13	10	11
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3	11	9	3	1	1	11	14	15	9	3	1	10	8	1	9	14	12	5	8
7	12	10	8	3	15	9	7	10	8	12	8	4	11	2	2	7	6	11	6
5	2	14	14	13	14	4	13	9	7	6	5	14	2	9	7	11	15	6	9
4	5	4	15	9	4	13	6	16	4	10	15	2	5	6	10	12	16	16	7
14	1	11	5	11	5	7	5	14	12	11	10	12	15	7	4	10	5	15	15
15	4	13	4	4	6	6	3	6	5	5	4	3	1	13	12	3	10	2	2
1	13	2	6	2	3	3	2	8	6	13	12	7	9	5	11	13	3	1	10
6	10	7	12	14	11	5	10	5	13	2	11	9	14	16	14	6	14	12	4
16	9	1	10	16	7	1	9	2	16	8	6	15	12	11	13	15	9	9	5
2	14	6	13	12	12	15	8	11	3	16	16	13	6	14	5	1	1	3	12
8	3	15	9	5	16	2	15	7	15	9	13	16	4	15	1	8	4	14	14
10	15	3	11	6	10	10	16	12	1	15	3	11	13	8	6	16	7	7	10
10	7	12	15	9	16	10	8	2	2	2	12	10	4	15	3	2	6	9	5
15	5	13	8	14	12	12	15	7	9	7	2	3	16	7	16	7	8	1	14
12	10	8	3	8	5	6	14	4	4	15	3	7	7	16	6	9	9	4	15
9	15	14	2	5	3	8	10	12	8	3	7	5	5	12	8	6	13	10	10
7	2	3	4	3	14	3	3	11	11	4	4	4	6	4	2	10	2	7	3
13	13	16	10	7	1	1	7	8	13	10	1	8	14	10	1	3	12	15	12
1	3	1	6	4	15	7	6	5	15	12	5	6	2	2	10	15	10	12	2
5	14	2	12	13	2	14	13	6	5	9	10	12	3	11	12	8	1	3	9
3	8	5	1	12	9	15	1	13	10	14	16	9	13	6	7	5	5	11	7
2	1	9	5	2	8	4	4	16	7	11	9	13	15	1	15	13	11	2	11
6	9	4	7	6	10	13	5	10	14	8	15	2	8	13	5	16	4	16	16
8	16	10	9	1	6	5	2	1	3	6	8	15	11	9	9	11	7	8	4
11	6	15	16	16	13	9	11	9	12	16	6	11	9	8	13	14	16	14	1
4	4	11	11	15	11	2	16	3	16	13	11	14	12	5	11	4	14	5	13
14	12	7	13	10	4	16	12	14	6	5	13	16	1	14	4	1	15	13	6
16	11	6	14	11	7	11	9	15	1	1	14	1	10	3	14	12	3	6	8

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

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2	9	16	16	16	2	6	4	9	10	3	2	15	10	2	3	11	13	10	1
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9	8	15	3	13	7	16	11	6	4	10	6	4	12	3	1	15	5	13	13
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1	11	6	5	2	13	12	7	4	14	14	13	10	2	16	15	6	10	5	16
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TABLE 15.7 PERMUTATIONS OF 16 (Continued)

9	2	4	7	11	8	9	12	11	3	10	8	11	14	16	2	13	2	2	15
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13	7	2	2	12	13	6	6	2	6	11	9	8	2	3	6	3	5	3	5
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TABLE 15.7 PERMUTATIONS OF 16 (Continued)

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4	2	6	6	15	8	10	15	13	9	8	9	12	13	10	3	16	8	9	8
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14	15	8	14	9	14	16	6	1	10	14	5	7	12	2	6	10	2	11	1
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15	14	1	10	12	10	10	16	6	4	8	11	1	9	3	14	8	3	7	3
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14	3	8	15	16	9	7	4	14	13	8	15	9	3	9	5	14	10	13	8
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9	8	2	4	7	14	15	11	13	8	11	14	3	12	3	16	13	5	16	2
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12	9	4	2	14	7	5	9	2	15	10	4	15	11	13	9	15	13	15	13
3	1	3	6	4	13	16	5	5	16	5	9	13	14	4	13	3	4	14	4

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

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12	16	10	10	5	8	13	13	4	13	15	1	11	4	16	5	6	6	8	9
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TABLE 15.7 PERMUTATIONS OF 16 (Continued)

15	2	11	8	9	2	11	15	3	1	15	16	1	2	1	3	12	1	4	3
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14	10	7	1	10	9	1	7	10	14	5	4	10	13	7	7	9	7	3	10
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7	14	16	15	13	12	6	3	6	8	8	5	15	3	8	2	7	5	8	7
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13	1	10	16	2	7	8	16	15	15	14	14	11	10	12	11	4	4	14	1
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7	1	3	9	5	12	2	1	5	6	12	12	10	2	14	15	3	10	9	2
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15	6	9	14	11	2	2	16	4	11	16	14	9	5	2	8	3	7	7	1
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7	11	3	8	14	4	12	7	13	13	2	4	16	16	15	13	2	10	13	10
14	12	16	2	15	14	13	14	15	14	1	10	3	2	11	15	10	1	16	2
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TABLE 15.7 PERMUTATIONS OF 16 (Continued)

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9	6	15	6	10	8	6	14	1	5	15	9	4	8	11	9	12	11	6	9
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9	16	10	5	2	13	14	12	15	11	14	7	16	11	9	9	3	13	11	1
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12	7	1	12	1	8	12	14	11	10	16	16	1	12	11	13	11	14	9	12
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14	8	14	15	7	10	16	13	12	15	5	11	8	7	13	11	16	8	6	16
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6	3	8	9	14	9	13	2	5	6	10	2	6	15	10	10	13	15	5	8
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9	12	16	8	10	7	4	15	5	14	11	3	8	15	13	16	14	1	1	15
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2	7	8	9	5	6	13	13	4	2	5	13	13	16	7	2	9	9	15	5
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10	2	12	14	3	11	5	8	13	8	7	12	1	11	11	10	5	13	4	1
11	16	6	4	1	16	12	14	16	13	1	5	7	8	3	12	12	10	7	2
14	10	5	13	8	10	7	6	12	9	9	6	12	4	12	3	7	4	8	13
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4	9	2	6	11	12	11	2	9	6	6	1	4	9	2	9	16	7	14	8
8	4	7	10	6	8	10	4	8	12	8	14	6	3	9	14	1	5	6	16
15	5	15	16	15	5	16	3	7	1	3	10	5	14	6	15	3	15	11	9
13	8	10	11	9	13	14	12	1	5	3	6	5	14	16	6	6	6	9	11
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3	6	4	15	7	9	1	9	15	4	15	16	11	7	14	5	15	16	2	7
12	11	11	1	16	1	8	10	10	3	4	4	16	13	1	8	10	3	12	6
5	14	13	2	2	4	9	7	3	15	2	8	14	12	5	1	11	2	5	10

TABLE 15.7. PERMUTATIONS OF 16 (Continued)

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11	3	16	1	12	15	4	15	15	6	4	2	10	2	9	4	11	15	16	10
7	16	15	10	10	11	13	13	7	4	10	10	12	9	3	1	1	10	2	2
4	5	1	6	5	6	11	16	12	11	11	1	9	11	4	16	2	8	5	4
14	4	8	5	3	4	5	14	1	12	12	11	4	10	12	7	12	14	14	14
3	9	14	11	15	9	16	7	3	16	15	7	2	16	7	13	14	3	3	12
2	14	5	14	14	7	1	3	5	14	3	12	13	3	14	2	15	1	4	1
9	2	3	3	13	3	14	2	9	7	9	4	7	7	2	12	16	7	13	7
12	13	13	7	7	8	6	9	14	3	7	16	6	5	13	9	8	9	12	11
16	1	9	2	11	13	10	8	10	15	14	13	3	6	10	5	13	11	7	8
15	12	6	4	4	5	7	12	16	2	8	5	5	13	5	8	7	16	6	6
8	15	12	8	2	16	9	5	2	13	2	9	14	1	1	6	4	2	9	15
5	8	4	13	1	12	2	4	13	1	13	14	16	15	15	3	10	13	1	9
13	7	10	12	9	2	8	6	6	5	1	8	8	14	8	15	9	6	8	13
11	7	16	10	15	3	1	14	1	2	15	4	3	14	8	15	1	5	16	6
1	6	1	6	7	7	6	6	15	5	10	12	6	11	7	4	14	15	2	10
3	3	12	5	6	11	8	11	2	12	16	14	2	15	9	11	12	14	13	7
10	1	10	11	3	9	7	2	11	16	3	7	15	5	13	7	13	12	9	13
12	8	7	16	12	2	3	1	16	3	1	8	10	1	16	5	4	13	4	4
6	15	4	8	8	5	9	13	9	10	6	1	11	12	3	10	7	8	5	1
9	11	6	15	1	8	2	4	6	13	5	3	4	13	6	8	3	9	1	16
13	16	3	4	11	14	4	12	5	11	7	5	9	7	11	3	10	6	6	15
14	4	11	14	4	10	5	10	12	8	8	16	7	2	4	12	2	11	7	2
5	5	2	9	9	13	10	9	13	15	2	15	1	10	15	2	15	3	10	5
4	14	8	3	16	4	16	16	7	7	4	2	14	4	5	16	8	2	3	12
7	12	5	12	5	16	15	5	14	6	9	6	16	9	14	13	9	4	15	14
8	2	9	2	14	1	12	8	4	4	13	11	13	16	1	6	11	16	11	8
16	10	15	13	10	15	14	7	10	14	11	10	8	8	2	14	5	7	8	11
15	9	14	1	2	12	13	15	3	1	12	9	12	6	10	9	16	10	12	9
2	13	13	7	13	6	11	3	8	9	14	13	5	3	12	1	6	1	14	3
7	3	11	1	15	5	13	11	14	9	1	2	11	6	5	11	1	10	15	7
4	4	6	11	12	8	12	16	11	3	7	11	13	13	10	15	15	1	5	14
12	15	13	4	13	4	3	14	5	4	14	1	1	11	16	14	6	7	10	15
9	8	16	9	11	1	14	5	1	11	8	15	2	8	8	9	14	5	6	12
10	2	7	2	10	14	6	8	2	7	9	14	5	15	9	3	9	9	1	5
15	13	12	12	5	9	7	7	10	1	11	9	9	12	1	12	5	8	3	3
8	1	2	13	14	12	1	12	12	8	6	4	16	1	4	13	12	16	2	4
5	9	4	14	6	2	15	13	15	5	2	12	6	5	3	2	2	13	9	2
14	10	5	3	9	13	8	6	16	2	13	13	7	7	15	8	13	11	4	1
3	16	1	15	2	15	4	4	13	6	10	7	4	16	14	1	4	3	14	9
11	14	8	7	3	6	9	9	9	12	12	3	12	9	12	5	8	2	16	16
6	7	9	10	1	7	11	2	7	15	16	6	3	4	11	10	3	6	7	11
13	6	14	16	16	10	2	10	6	13	15	16	14	14	2	6	7	4	8	13
1	5	15	5	4	16	16	3	8	14	5	5	8	2	6	4	11	14	13	6
16	11	3	8	7	3	5	15	4	10	4	10	10	10	13	7	16	12	11	10
2	12	10	6	8	11	10	1	3	16	3	8	15	3	7	16	10	15	12	8
8	2	1	14	10	6	1	1	8	3	7	7	1	2	12	2	12	15	3	4
6	10	2	2	6	13	9	13	7	12	11	1	13	3	16	4	6	3	6	8
11	9	3	4	3	10	4	7	4	4	15	8	4	9	8	7	4	10	7	2
5	3	14	10	9	8	6	10	2	15	10	14	2	6	1	9	5	2	15	10
10	15	7	16	13	12	12	15	11	6	4	11	6	12	11	8	11	11	16	3
12	16	8	11	12	5	11	5	12	11	8	9	5	7	9	13	8	13	14	9
16	12	13	5	7	4	8	14	16	13	3	6	11	15	10	14	1	16	9	11
15	5	4	12	1	3	2	12	10	8	2	4	9	8	5	15	3	4	12	13
2	6	10	1	16	14	3	8	1	9	13	2	14	10	7	5	16	8	10	1
3	4	6	8	5	16	13	2	6	7	5	5	15	13	14	6	10	7	2	12
9	11	16	13	8	1	15	4	14	1	14	10	16	5	6	10	2	1	11	15
1	13	9	7	4	9	10	16	5	16	1	12	3	1	3	3	7	5	8	6
4	1	11	15	15	11	14	6	9	5	16	13	10	14	13	11	15	14	4	14
13	8	12	9	2	2	16	9	13	10	9	15	8	16	4	12	13	6	13	5
14	14	5	3	11	7	5	11	15	2	6	16	12	11	15	1	9	12	5	7
7	7	15	6	14	15	7	3	3	14	12	3	7	4	2	16	14	9	1	16

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

13	1	9	7	7	13	6	15	1	12	11	10	11	7	5	3	9	11	10	3
2	12	15	4	16	10	2	16	16	3	13	12	1	3	15	1	15	1	7	5
3	9	2	9	9	1	5	10	9	13	7	15	13	12	4	11	16	12	11	10
5	2	4	10	12	8	13	6	2	6	9	9	15	13	1	5	12	4	5	1
4	6	7	8	13	9	15	4	5	5	14	5	4	4	2	8	10	14	8	14
14	4	8	1	11	7	12	14	4	7	3	16	12	15	10	4	5	3	4	16
1	14	13	12	5	11	10	5	14	2	10	3	5	2	7	12	13	15	15	8
16	11	6	5	3	14	8	1	12	8	5	14	8	5	3	15	1	10	14	13
15	15	12	15	10	16	3	11	10	1	15	8	14	8	12	10	11	13	3	9
8	7	10	16	8	12	4	2	7	10	4	2	7	6	16	2	4	6	2	11
7	16	5	14	15	6	1	3	15	14	2	11	3	11	6	7	6	8	13	12
6	10	14	13	14	5	14	12	13	9	12	7	9	14	13	9	8	5	1	6
11	8	11	11	4	15	7	13	3	16	8	4	2	1	9	6	14	7	9	7
12	13	3	2	1	3	16	7	8	4	6	13	16	9	14	14	3	2	12	4
9	5	16	6	2	2	9	8	11	15	1	6	6	16	11	16	7	9	16	2
10	3	1	3	6	4	11	9	6	11	16	1	10	10	8	13	2	16	6	15
10	4	1	16	14	11	1	11	8	11	16	15	9	4	2	3	10	10	16	5
13	2	5	8	6	10	3	15	12	14	10	3	16	1	9	16	8	1	3	9
9	16	15	3	11	7	11	1	14	9	12	7	1	16	1	2	1	7	2	3
8	14	11	6	15	12	6	10	5	1	3	1	11	13	4	5	7	14	7	10
3	5	8	11	2	9	15	4	10	5	8	13	14	12	11	11	3	16	13	13
11	3	4	12	8	13	9	9	1	16	7	16	3	8	15	8	4	8	15	16
14	11	12	13	13	14	16	12	3	4	4	9	5	2	5	13	9	9	11	14
5	9	7	1	5	1	13	14	2	15	2	2	2	11	16	6	6	15	5	12
1	8	6	2	4	2	10	6	4	8	15	12	15	7	13	15	11	3	9	15
6	7	9	4	10	3	8	8	15	7	14	11	6	15	14	2	4	6	6	8
4	6	2	5	7	8	4	7	13	13	9	14	10	10	7	9	15	12	1	4
12	12	14	10	9	5	7	5	7	10	5	10	8	9	6	7	5	2	12	1
15	1	10	7	16	16	5	13	9	6	1	6	7	5	3	10	13	13	8	6
7	10	3	9	3	4	2	2	16	12	6	4	4	14	10	14	14	5	10	11
16	15	13	15	1	6	12	16	11	3	11	8	12	3	8	1	12	11	4	7
2	13	16	14	12	15	14	3	6	2	13	5	13	6	12	12	16	4	14	2
15	15	8	2	4	10	16	12	12	9	3	6	11	10	16	8	7	4	9	1
5	6	1	4	15	15	9	2	1	16	10	11	13	4	9	4	15	1	16	16
12	8	3	1	1	2	8	3	6	4	12	15	15	6	5	7	6	16	2	2
1	12	15	16	2	14	10	7	15	3	7	1	4	15	14	3	9	5	14	13
2	16	5	12	11	4	12	13	10	14	16	7	2	9	6	1	11	6	6	14
16	11	7	10	12	16	6	9	14	13	5	9	3	5	13	9	4	7	4	6
8	2	9	6	8	3	4	16	7	7	8	14	14	11	12	5	10	12	15	10
6	4	13	5	14	5	11	11	5	6	13	12	1	3	3	2	1	3	10	15
13	14	14	8	9	6	3	8	2	10	1	2	6	14	1	12	13	15	3	4
10	1	2	9	10	1	15	1	13	11	6	8	16	1	2	16	2	10	1	5
7	9	6	3	5	9	14	4	11	1	14	4	12	13	11	14	12	9	13	7
9	7	16	7	6	13	5	5	3	12	2	5	8	8	4	13	16	14	5	12
14	3	12	13	13	7	13	10	8	8	11	3	10	2	8	11	8	11	8	8
4	5	11	15	16	11	2	15	16	5	4	16	9	12	10	15	5	8	7	11
11	10	10	11	3	8	7	14	9	15	15	10	5	7	7	10	14	13	12	3
3	13	4	14	7	12	1	6	4	2	9	13	7	16	15	6	3	2	11	9
11	1	3	7	8	10	11	8	15	9	15	8	13	8	4	8	4	3	6	15
7	8	2	9	14	4	7	10	3	7	16	7	2	10	7	5	2	16	14	8
1	15	11	3	7	9	12	13	4	13	4	3	10	14	5	2	3	5	11	11
15	13	14	16	4	14	15	16	10	14	13	6	8	13	11	12	7	10	4	7
12	11	1	8	5	11	9	4	5	16	14	1	9	1	13	1	14	11	2	9
4	9	4	15	15	7	3	6	8	11	12	11	15	16	12	3	12	9	13	1
9	6	12	5	13	13	4	11	12	15	3	15	3	4	10	6	5	1	7	4
6	14	6	11	12	5	6	1	14	1	1	12	12	2	3	7	8	13	16	16
10	12	7	4	16	1	13	9	1	12	8	9	1	11	2	9	15	14	3	12
13	16	16	10	1	15	16	7	11	10	10	13	14	15	15	10	10	8	9	2
14	5	9	12	3	16	8	15	2	5	7	4	16	12	14	4	11	4	12	14
3	2	8	2	11	8	1	12	9	3	5	2	11	5	16	15	9	12	5	13
16	7	13	13	10	6	10	14	6	4	9	14	7	6	9	13	1	2	1	6
8	4	5	1	6	3	2	5	16	8	6	16	4	7	1	16	16	7	15	3
2	10	15	14	9	12	14	3	7	2	2	10	6	3	6	11	6	6	8	5
5	3	10	6	2	2	5	2	13	6	11	5	5	9	8	14	13	15	10	10

TABLE 15.7 PERMUTATIONS OF 16 (Continued)

10	14	13	1	6	6	13	1	8	4	9	9	1	2	9	3	1	15	12	8
7	12	2	5	2	16	2	14	16	16	14	10	15	3	14	15	9	8	15	10
11	6	10	10	10	8	1	9	3	14	12	11	7	5	1	5	10	12	11	5
14	11	16	16	15	12	3	12	9	1	16	8	11	11	0	7	5	3	10	15
5	16	5	8	11	13	16	7	5	10	11	1	3	14	4	1	15	1	4	16
15	15	8	14	1	1	4	8	7	8	3	15	14	6	2	14	6	11	7	4
13	7	14	9	14	11	14	2	15	5	10	7	4	8	7	10	4	6	5	9
4	2	1	11	13	14	15	16	4	13	1	2	9	13	5	4	13	9	9	13
3	4	3	12	8	10	11	3	10	11	5	12	10	7	3	16	7	13	1	14
12	9	6	13	12	5	9	15	12	6	2	16	5	10	15	2	11	2	8	7
16	3	11	6	4	9	7	6	11	3	15	4	8	1	16	11	14	4	2	6
8	13	12	4	3	7	12	13	13	7	8	6	6	4	11	13	12	10	14	2
1	10	4	15	5	2	6	11	2	9	6	13	13	16	8	8	8	14	16	12
6	1	9	7	7	4	8	5	1	12	4	3	12	12	13	12	16	16	13	11
9	5	15	3	9	15	10	4	6	15	7	5	2	9	10	6	2	7	6	3
2	8	7	2	10	3	5	10	14	2	13	14	16	15	12	9	3	5	3	1
7	11	1	15	10	2	14	4	14	14	6	4	15	4	6	11	11	5	13	10
2	6	14	8	12	16	6	10	11	7	5	12	1	7	14	9	15	12	7	14
12	8	5	14	1	7	11	16	5	12	16	10	7	9	9	3	2	6	5	7
11	2	9	4	2	6	5	1	9	11	7	5	9	2	8	13	1	10	14	5
6	4	6	3	5	5	10	15	3	13	9	15	12	10	2	10	12	15	16	16
16	14	7	13	9	1	16	13	12	10	3	9	10	14	7	7	3	7	2	12
5	7	15	12	3	3	4	5	6	9	11	13	8	13	16	2	6	14	8	11
8	15	4	2	16	8	8	3	10	8	1	2	13	12	5	8	10	3	3	13
9	5	10	11	8	13	13	14	4	16	13	1	3	6	3	4	14	9	4	2
1	16	12	10	6	9	3	7	16	1	12	8	14	1	10	1	4	8	11	3
3	9	8	1	4	14	1	12	7	4	10	3	5	11	11	14	13	16	10	9
14	1	11	7	7	4	9	8	8	3	4	14	4	5	12	16	5	1	6	1
10	12	2	5	11	12	15	11	13	2	14	11	16	8	1	12	7	13	12	4
4	10	13	16	14	11	7	6	2	6	15	6	11	15	15	6	9	11	15	8
13	3	3	9	13	15	2	9	15	15	8	16	6	3	4	15	16	4	1	15
15	13	16	6	15	10	12	2	1	5	2	7	2	16	13	5	8	2	9	6

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